# WARF0235.ST25.txt SEQUENCE LISTING

<110>	Helge Austi Naess Song, Jiang Brade Buell	n-Ph , Sa Jun , Ji en,	nill: ara H nqi iminq Jame	ips, Krist g es Ma	Sand tine										
<120>	POTAT	O GE	ENES	FOR	RES	ISTAI	NCE '	ro L	ATE I	BLIG	нт				
<130>	WARF-	0235	5												
<150> <151>	US 60 2003-			5											
<160>	125														
<170>	Paten	tIn	vers	sion	3.2										
<210> <211> <212> <213>	1 2940 DNA Solan	um b	oulbo	ocast	tanur	n									
<220> <221> <222>	CDS (1)	(294	10)												
<400> atg gc Met Ala 1	1 t gaa ( a Glu <i>i</i>	gct Ala	ttc Phe 5	att Ile	caa Gln	gtt Val	gtg Val	cta Leu 10	gac Asp	aat Asn	ctc Leu	act Thr	tct Ser 15	ttc Phe	48
ctc aad Leu Ly	s Gly (	gaa Glu 20	ctt Leu	gta Val	ttg Leu	ctt Leu	ttc Phe 25	ggt Gly	ttt Phe	caa Gln	gat Asp	gag Glu 30	ttc Phe	caa Gln	96
agg ct															144
caa gad Gln Gl 50	g aag ( u Lys (	caa Gln	ctc Leu	aac Asn	gac Asp 55	aag Lys	cct Pro	cta Leu	gaa Glu	aat Asn 60	tgg Trp	ttg Leu	caa Gln	aaa Lys	192
ctc aar Leu Asi 65					Glu	Val	Asp		Ile	Leu	Asp	Glu		Lys	240
act aad Thr Ly															288
aag gt: Lys Va	l Ile :														336
atg aaa Met Ly:															384
caa gaa Gln Gli 13	u Lys				_		-	_			_				432
gtg tta Val Le															480

160

145					150					155					160	
			atc Ile													528
gtc Val	ctc Leu	cca Pro	ata Ile 180	ctt Leu	ggt Gly	atg Met	Gly ggg	gga Gly 185	cta Leu	gga Gly	aag Lys	acg Thr	act Thr 190	ctt Leu	tcc Ser	576
caa Gln	atg Met	gtc Val 195	ttc Phe	aat Asn	gat Asp	cag Gln	aga Arg 200	gta Val	act Thr	gag Glu	cgt Arg	ttc Phe 205	tat Tyr	ccc Pro	aaa Lys	624
ata Ile	tgg Trp 210	att Ile	tgc Cys	atc Ile	tcg Ser	gat Asp 215	gat Asp	ttt Phe	aat Asn	gag Glu	aag Lys 220	agg Arg	ttg Leu	ata Ile	aag Lys	672
gca Ala 225	ata Ile	gta Val	gaa Glu	tct Ser	att Ile 230	gaa Glu	gly Gly	aag Lys	tcc Ser	ctc Leu 235	agt Ser	gac Asp	atg Met	gac Asp	ttg Leu 240	720
			caa Gln													768
ttc Phe	ctt Leu	gtc Val	tta Leu 260	gat Asp	gat Asp	gtt Val	tgg Trp	aat Asn 265	gaa Glu	gat Asp	caa Gln	cat His	aag Lys 270	tgg Trp	gct Ala	816
			gca Ala													864
			cgt Arg													912
			ttg Leu													960
			gca Ala													1008
			aag Lys 340													1056
			ctt Leu													1104
			gtg Val													1152
			ctg Leu													1200
			caa Gln													1248
			aag Lys 420													1296

								WAR	F023	5.ST	'25.t	xt			
					ttg Leu										1344
					agg Arg 455										1392
					aag Lys	_		_				_	_	_	1440
	_			-	aac Asn			_	_			_	_		1488
-			-		tat Tyr	_	_	-				_	-		153.6
					tca Ser										1584
			_		tcg Ser 535										1632
					aga Arg										1680
					aga Arg										1728
-				-	gac Asp				_	_					1776
					cga Arg										1824
					ata Ile 615										1872
					aag Lys										1920
					ggc Gly										1968
					gca Ala										2016
_				_	ctg Leu	-		_		-				_	2064
_		-	_		gaa Glu 695	-							_		2112
	-				ttc Phe				_			-		_	2160

705					710	WARF0235.ST25.txt 710 715										
			gtt Val													2208
-		_	tca Ser 740	_							_		_			2256
			tta Leu													2304
aat Asn	gtt Val 770	cat His	cct Pro	gga Gly	agg Arg	ttt Phe 775	cca Pro	tcc Ser	ttg Leu	agg Arg	aaa Lys 780	ctt Leu	gtt Val	ata Ile	tgg Trp	2352
			aat Asn													2400
			ctt Leu													2448
			ctt Leu 820													2496
			ttg Leu													2544
			gat Asp													2592
			gca Ala													2640
			ttg Leu													2688
			gaa Glu 900													2736
			tta Leu													2784
			tgt Cys													2832
			att Ile			-			-		-		_			2880
			gaa Glu	-					-					_		2928

2940

cta tat gag tga Leu Tyr Glu

- <210> 2 <211> 979
- <212> PRT
- <213> Solanum bulbocastanum
- <221> misc\_feature
- <222> (119)...(119)
- <223> The 'Xaa' at location 119 stands for Ile, or Phe.
- <400> 2
- Met Ala Glu Ala Phe Ile Gln Val Val Leu Asp Asn Leu Thr Ser Phe
- Leu Lys Gly Glu Leu Val Leu Phe Gly Phe Gln Asp Glu Phe Gln
- Arg Leu Ser Ser Met Phe Ser Thr Ile Gln Ala Val Leu Glu Asp Ala 40
- Gln Glu Lys Gln Leu Asn Asp Lys Pro Leu Glu Asn Trp Leu Gln Lys
- Leu Asn Ala Ala Thr Tyr Glu Val Asp Asp Ile Leu Asp Glu Tyr Lys
- Thr Lys Ala Thr Arg Phe Leu Gln Ser Glu Tyr Gly Arg Tyr His Pro
- Lys Val Ile Pro Phe Arg His Lys Val Gly Lys Arg Met Asp Gln Val
- Met Lys Lys Leu Asn Ala Xaa Ala Glu Glu Arg Lys Lys Phe His Leu
- Gln Glu Lys Ile Ile Glu Arg Gln Ala Ala Thr Arg Glu Thr Gly Ser
- Val Leu Thr Glu Pro Gln Val Tyr Gly Arg Asp Lys Glu Lys Asp Glu
- Ile Val Lys Ile Leu Ile Asn Thr Ala Ser Asp Ala Gln Lys Leu Ser
- Val Leu Pro Ile Leu Gly Met Gly Gly Leu Gly Lys Thr Thr Leu Ser
- Gln Met Val Phe Asn Asp Gln Arg Val Thr Glu Arg Phe Tyr Pro Lys 200
- Ile Trp Ile Cys Ile Ser Asp Asp Phe Asn Glu Lys Arg Leu Ile Lys 215
- Ala Ile Val Glu Ser Ile Glu Gly Lys Ser Leu Ser Asp Met Asp Leu Page 5

230

Ala	Pro	Leu	Gln	Lys 245	Lys	Leu	Gln	Glu	Leu 250	Leu	Asn	Gly	Lys	Arg 255	Tyr
Phe	Leu	Val	Leu 260	Asp	Asp	Val	Trp	Asn 265	Glu	Asp	Gln	His	Lys 270	Trp	Ala
Asn	Leu	Arg 275	Ala	Val	Leu	Lys	Val 280	Gly	Ala	Ser	Gly	Ala 285	Phe	Val	Leu
Thr	Thr 290	Thr	Arg	Leu	Glu	Lys 295	Val	Gly	Ser	Ile	Met 300	Gly	Thr	Leu	Gln
Pro 305	Tyr	Glu	Leu	Ser	Asn 310	Leu	Ser	Pro	Glů	Asp 315	Cys	Trp	Phe	Leu	Phe 320
Met	Gln	Arg	Ala	Phe 325	Gly	His	Gln	Glu	Glu 330	Ile	Asn	Pro	Asn	Leu 335	Met
Ala	Ile	Gly	Lys 340	Glu	Ile	Val	Lys	Lys 345	Cys	Gly	Gly	Val	Pro 350	Leu	Ala
Ala	Lys	Thr 355	Leu	Gly	Gly	Ile	Leu 360	Arg	Phe	Lys	Arg	Glu 365	Glu	Arg	Glu
Trp	Glu 370	His	Val	Arg	Asp	Ser 375	Pro	Ile	Trp	Asn	Leu 380	Pro	Gln	Asp	Glu
Ser 385	Ser	Ile	Leu	Pro	Ala 390	Leu	Arg	Leu	Ser	Tyr 395	His	His	Leu	Pro	Leu 400
Asp	Leu	Arg	Gln	Cys 405	Phe	Val	Tyr	Cys	Ala 410	Val	Phe	Pro	Lys	Asp 415	Thr
Lys	Met	Ala	Lys 420	Glu	Asn	Leu	Ile	Ala 425	Phe	Trp	Met	Ala	His 430	Gly	Phe
Leu	Leu	Ser 435	Lys	Gly	Asn	Leu	Glu 440	Leu	Glu	Asp	Val	Gly 445	Asn	Glu	Val
Trp	Asn 450	Glu	Leu	Tyr	Leu	Arg 455	Ser	Phe	Phe	Gln	Glu 460	Ile	Glu	Val	Glu
Ser 465	Gly	Lys	Thr	Tyr	Phe 470	Lys	Met	His	Asp	Leu 475	Ile	His	Asp	Leu	Ala 480
Thr	Ser	Leu	Phe	Ser 485	Ala	Asn	Thr	Ser	Ser 490	Ser	Asn	Ile	Arg	Glu 495	Ile
Asn	Ala	Asn	Tyr 500	Asp	Gly	Tyr	Met	Met 505	Ser	Ile	Gly	Phe	Ala 510	Glu	Val

Val	Ser	Ser 515	Tyr	Ser	Pro	Ser	Leu 520	Leu	Gln	Lys	Phe	Val 525	Ser	Leu	Arg
Val	Leu 530	Asn	Leu	Arg	Asn	Ser 535	Asn	Leu	Asn	Gln	Leu 540	Pro	Ser	Ser	Ile
Gly 545	Asp	Leu	Val	His	Leu 550	Arg	Tyr	Leu	Asp	Leu 555	Ser	Gly	Asn	Phe	Arg 560
Ile	Arg	Asn	Leu	Pro 565	Lys	Arg	Leu	Cys	Lys 570	Leu	Gln	Asn	Leu	Gln 575	Thr
Leu	Asp	Leu	His 580	Tyr	Cys	Asp	Ser	Leu 585	Ser	Cys	Leu	Pro	Lys 590	Gln	Thr
Ser	Lys	Leu 595	Gly	Ser	Leu	Arg	Asn 600	Leu	Leu	Leu	Asp	Gly 605	Cys	Ser	Leu
Thr	Ser 610	Thr	Pro	Pro	Arg	Ile 615	Gly	Leu	Leu	Thr	Cys 620	Leu	Lys	Ser	Leu
Ser 625	Cys	Phe	Val	Ile	Gly 630	Lys	Arg	Lys	Gly	His 635	Gln	Leu	Gly	Glu	Leu 640
Lys	Asn	Leu	Asn	Leu 645	Tyr	Gly	Ser	Ile	Ser 650	Ile	Thr	Lys	Leu	Asp 655	Arg
Val	Lys	Lys	Asp 660	Thr	Asp	Ala	Lys	Glu 665	Ala	Asn	Leu	Ser	Ala 670	Lys	Ala
Asn	Leu	His 675	Ser	Leu	Cys	Leu	Ser 680	Trp	Asp	Leu	Asp	Gly 685	Lys	His	Arg
Tyr	Asp 690	Ser	Glu	Val	Leu	Glu 695	Ala	Leu	Lys	Pro	His 700	Ser	Asn	Leu	Lys
Tyr 705	Leu	Glu	Ile	Asn	Gly 710	Phe	Gly	Gly	Ile	Arg 715	Leu	Pro	Asp	Trp	Met 720
Asn	Gln	Ser	Val	Leu 725	Lys	Asn	Val	Val	Ser 730	Ile	Arg	Ile	Arg	Gly 735	Cys
Glu	Asn	Cys	Ser 740	Cys	Leu	Pro	Pro	Phe 745	Gly	Glu	Leu	Pro	Cys 750	Leu	Glu
Ser	Leu	Glu 755	Leu	His	Thr	Gly	Ser 760	Ala	Asp	Val	Glu	Туг 765	Val	Glu	Asp
Asn	Val 770	His	Pro	Gly	Arg	Phe 775	Pro	Ser	Leu	Arg	Lys 780	Leu	Val	Ile	Trp
Asp	Phe	Ser	Asn	Leu	Lys	Gly	Leu	Leu	Lys		Glu age '	_	Glu	Lys	Gln

WARF0235.ST25.txt 785 790 Phe Pro Val Leu Glu Glu Met Thr Phe Tyr Trp Cys Pro Met Phe Val Ile Pro Thr Leu Ser Ser Val Lys Thr Leu Lys Val Ile Val Thr Asp Ala Thr Val Leu Arg Ser Ile Ser Asn Leu Arg Ala Leu Thr Ser Leu 840 Asp Ile Ser Asp Asn Val Glu Ala Thr Ser Leu Pro Glu Glu Met Phe 860 Lys Ser Leu Ala Asn Leu Lys Tyr Leu Lys Ile Ser Phe Phe Arg Asn Leu Lys Glu Leu Pro Thr Ser Leu Ala Ser Leu Asn Ala Leu Lys Ser 890 Leu Lys Phe Glu Phe Cys Asp Ala Leu Glu Ser Leu Pro Glu Glu Gly 905

Val Lys Gly Leu Thr Ser Leu Thr Glu Leu Ser Val Ser Asn Cys Met

Met Leu Lys Cys Leu Pro Glu Gly Leu Gln His Leu Thr Ala Leu Thr 935

Thr Leu Thr Ile Thr Gln Cys Pro Ile Val Phe Lys Arg Cys Glu Arg

Gly Ile Gly Glu Asp Trp His Lys Ile Ala His Ile Pro Tyr Leu Thr 965 970

Leu Tyr Glu

<210> 3 <211> 8569 <212> DNA

<213> Solanum bulbocastanum

<220>

<221> exon (2345)..(2804) <222>

<220>

<221> Intron

<222> (2805)..(3456)

<220>

<221> exon

<222> (3457)..(5836)

<400> 3

WARF0235.ST25.txt cgggatcctg tcacataaat tgacacaaag ggagtacttg ttaatgttgt aattattggc 60 gaacaataat gttgttgatt atcactttct gaataagtgt tgtgtcactt ggaaaaaaca 120 ccaaatagaa ctattcatgt tttttcttta gtatatataa atatgatctt taacttaatt 180 tcagcagaca gtcatgatct ttaactttaa atgtgcacaa gtagattgac aggcttgcta 240 attgagtgtc tgttataatc agtattaaat actctcaagg taatagtata ttccagacaa 300 attttgtgtt accaaattaa atatatttct aaaactctcc tcaaagtagt taatatactt 360 ttgagtgttg tatcatgttt ttaatataaa atgttaaaat ttagatgaaa tttactttct 420 agttaaattg gtcaaagttg aaagaatttc aagtgaaaaa gtttttaata atttgacttt 480 tatgctatat ttttttaaag ttgaacgact ttttaataaa aaagaataat aaaattatat 540 gataattttt ataatacaat ggcctttata tgatgaaaaa aaaagaaaga aattagatga 600 caacaatgtc caaaaataat cttaaagaat tacgatttat atataataaa attaaattta 660 aaatttgatg aaaaaataga gaaaagagga agatgatgaa gtgaaatgac gtggtggtgg 720 gtccatgtga cataaaaaaa aattctctta aataatcctt tcatactaat gataaatttt 780 ttttttttt ttttttac taattgcgta tagagaaaag gaaaatgggg cggtaattac 840 aaagtaggga atcgaacttt atcaacaagt tgagagttca agtaatcaac caactaaact 900 actaaaattt ttctaattaa tgataattgt aattcattta gcataaaaaa tttcattgca 960 cttactttta gagttttgaa aacagtactt catctattct atattaatta aattttctat 1020 attaattaaa tttgtgaggt aatacaaact tattaagaaa aatatttaag gacataattt 1080 aactcatatt tttcactatt gttttttgtg aaatcataaa tataactttg taaatagtgc 1140 aatttatctc ctagaagcaa atttcaccaa agaaaagggc aaagatggaa aagaaactaa 1200 atattcatct taaactttga acaattcaat tattttgaac aatgaaaaaa atctcaaaaa 1260 ttcaattaat atgaaatgga gagagtaact ttattttaga ggcaaaaaat tagtactcca 1320 tccgttcact tttatttgtc atgttgcgct tttcgaaagt caatttgact aatttttaaa 1380 gctaaattag attacactaa ttcaatattt taaacagaaa aattagatat tcaaaaacta 1440 tacaaaaaat attatacatt gcaatttttt gcatatcaat atgataaaaa aatatattgt 1500 aaaatattag tcaaaatttt tatagtttga ctctaatcat gaaaagtata ataattaata 1560 gtggacggag gaagtattgt ctttccagat ttgtggccat ttttgggcca agggccatta 1620 gcagttctct tcattttcta cttctgtctc atattagatg ggcatcttac taaaaatatt 1680 tgtctcatat tacttgatta tttattaaat caaaaagaat taattaattt tttctcattt 1740 tacccctaca attaatatag ttttaaaaagt tttaaacaaa ttttgaagaa tcaaaatttc 1800 tttttgcaag agacttatta atataaacaa aggataaaaat aataaaattt gtcaatttat 1860 tgacgatcac ttaataatca tataaaatag aatatgttta tctaatatga gacggagaaa 1920 atatatccta aaatattttt ggacagatat gtgatattct aaccattcac tagactatat 1980 tatgcatttt agccgccaat gacttatttc agctttaatt aattaggaaa gaggaaactg 2040 ccaatgagga agagtagggg cgtagttgct gtcgacgaaa aaaagataat actcactctt 2100

WARF0235.ST25.txt	
ttcgattttt attttattt atcactttta acctatcatg taaaaagata attattttt	2160
tcatgcttta tccttagtat taaacaattt aatagggatt attttgtaaa atatttatat	2220
gaataattgt tttcgtaatg aatttgtccg gtcaaacaat gataaataaa aacgaatgaa	2280
gagagtagaa aacaaaacaa aagaacaagt tgacaacttg agagattaaa agggtccaaa	2340
acgc ctt gga ttt tga gat tcc ata tgt gaa att tcc atg aaa taa ttg Leu Gly Phe Asp Ser Ile Cys Glu Ile Ser Met Lys Leu 1 5 10	2389
aat ttg tat tat tac aag tca aac ttc cca ttt cat tcc aac tag cca Asn Leu Tyr Tyr Tyr Lys Ser Asn Phe Pro Phe His Ser Asn Pro 15 20 25	2437
tct tgg ttt caa aat tac aca ttc att cat tca cag atc taa tat tct Ser Trp Phe Gln Asn Tyr Thr Phe Ile His Ser Gln Ile Tyr Ser 30 35 40	2485
taa tag tga ttt cca cat atg gct gaa gct ttc att caa gtt ctg cta Phe Pro His Met Ala Glu Ala Phe Ile Gln Val Leu Leu 45 50 55	2533
gac aat ctc act tct ttc ctc aaa ggg gaa ctt gta ttg ctt ttc ggt Asp Asn Leu Thr Ser Phe Leu Lys Gly Glu Leu Val Leu Phe Gly 60 65 70	2581
ttt caa gat gag ttc caa agg ctt tca agc atg ttt tct aca att caa Phe Gln Asp Glu Phe Gln Arg Leu Ser Ser Met Phe Ser Thr Ile Gln 75 80 85	2629
gcc gtc ctt gaa gat gct cag gag aag caa ctc aac aac aag cct cta Ala Val Leu Glu Asp Ala Gln Glu Lys Gln Leu Asn Asn Lys Pro Leu 90 95 100	2677
gaa aat tgg ttg caa aaa ctc aat gct gct aca tat gaa gtc gat gac Glu Asn Trp Leu Gln Lys Leu Asn Ala Ala Thr Tyr Glu Val Asp Asp 105 110 115 120	2725
atc ttg gat gaa tat aaa acc aag gcc aca aga ttc tcc cag tct gaa Ile Leu Asp Glu Tyr Lys Thr Lys Ala Thr Arg Phe Ser Gln Ser Glu 125 130 135	2773
tat ggc cgt tat cat cca aag gtt atc cct t tccgtcacaa ggtcgggaaa Tyr Gly Arg Tyr His Pro Lys Val Ile Pro 140 145	2824
aggatggacc aagtgatgaa aaaactaaag gcaattgctg aggaaagaaa gaattttcat	2884
ttgcacgaaa aaattgtaga gagacaagct gttagacggg aaacaggtac tcatcttaaa	2944
ttagaattac aacaactaag tttatattca tttttttggc aattatgaaa ttcagaaaag	3004
ggttaaatat actcatgtcc tatcgtaaat agtgtaaata tacctctcgt tgtactttcg	3064
atctgaatat acttgtcaaa tctggcaagc tcagaatcaa attatccacc ccaactttta	3124
aatactcgat atctttagaa atccacctgt ctaactcatc cactacccat tccctttgct	3184
ttgaattett ttetttaeet ataaaegtgg aacaetegat eegttttget tttettaaca	3244
aagcagctca gagaaaagag gttttcttct attctgtttc tctgtgtgct gcacttgggt	3304
ccttaatccc attaaaaaca gggcatgtta atcccaacga cggtagcctt tcctgacagc	3364
tgactgtaaa ttttgtctaa caaagaaaaa aaaagattag acatgttttt ccttgtcatt	3424
gattaggctg gatttctttc agagtggaac at ag ggg ata tat tgg acc aaa Gly Ile Tyr Trp Thr Lys	3476

					tat Tyr					tga	tag		agg Arg 165	-		3524
					ctc Leu											3572
					atg Met											3620
					tat Tyr 205											3668
					aat Asn	_	_	_	-					-		3716
					ggg Gly											3764
					aga Arg											3812
					gat Asp											3860
					gga Gly 285							_	_	_	-	3908
					ctt Leu											3956.
					gtt Val											4004
					aag Lys											4052
					aag Lys											4100
					ctg Leu 365											4148
caa Gln	cgt Arg	gca Ala	ttt Phe	gga Gly 380	cac His	caa Gln	gaa Glu	gaa Glu	ata Ile 385	aat Asn	cca Pro	aac Asn	ctt Leu	gtg Val 390	gca Ala	4196
		_			gtg Val			_						-	-	4244
					att Ile											4292

	cat His 425								aat	ttg	cct		gat			4340
	att Ile															4388
ttg Leu	aaa Lys	caa Gln	tgc Cys	ttt Phe 460	gcg Ala	tat Tyr	tgt Cys	gcg Ala	gtg Val 465	ttc Phe	cca Pro	aag Lys	gat Asp	gcc Ala 470	aaa Lys	4436
	aaa Lys															4484
tta Leu	tca Ser	aaa Lys 490	gga Gly	aac Asn	atg Met	gag Glu	cta Leu 495	gag Glu	gat Asp	gtg Val	ggc Gly	gat Asp 500	gaa Glu	gta Val	tgg Trp	4532
	gaa Glu 505															4580
ggt Gly 520	aaa Lys	act Thr	tat Tyr	ttc Phe	aag Lys 525	atg Met	cat His	gat Asp	ctc Leu	atc Ile 530	cat His	gat Asp	ttg Leu	gca Ala	aca Thr 535	4628
	ctg Leu															4676
aaa Lys	cac His	agt Ser	tac Tyr 555	aca Thr	cat His	atg Met	atg Met	tcc Ser 560	att Ile	ggt Gly	ttc Phe	gcc Ala	gaa Glu 565	gtg Val	gtg Val	4724
	ttt Phe															4772
	aat Asn 585															4820
	cta Leu															4868
Ser	ctt Leu	Pro	Lys	Gln 620	Leu	Cys	Lys	Leu	Gln 625	Asn	Leū	Gln	Thr	Leu 630	Asp	4916
	caa Gln															4964
	ggt Gly	-		-					-		-	-		-		5012
_	atg Met 665							_				_				5060
	ttt Phe															5108

	_	_					
					tct gca aa Ser Ala Ly 72	's Gly Asn	5204
	er Leu Se				gga cca ca Gly Pro Hi 740		5252
			Leu Glu		aaa cca ca Lys Pro Hi 755		5300
					atc cat ct Ile His Le		5348
		r Val Leu			tct att ct Ser Ile Le		5396
					ggt gat ct Gly Asp Le 80	eu Pro Cys	5444
	er Leu Gl				gat gtg ga Asp Val Gl 820		5492
	•		His Ser		ccc aca ag Pro Thr Ar 835		5540
					ttt ggt ag Phe Gly Se		5588
gga ttg ct Gly Leu Le	g aaa aa eu Lys Ly 86	s Glu Gly	gaa gag Glu Glu	caa ttc Gln Phe 865	cct gtg ct Pro Val Le	t gaa gag u Glu Glu 870	5636
					tct tct aa Ser Ser As 88	n Leu Arg	5684
	r Ser Le				gta gct ac Val Ala Th 900		5732
cca gaa ga Pro Glu Gl 905	ig atg tto u Met Pho	c aaa aac e Lys Asn 910	Leu Ala	aat ctc Asn Leu	aaa tac tt Lys Tyr Le 915	g aca atc u Thr Ile	5780
					agc ttg gc Ser Leu Al		5828
aat gct tt Asn Ala	gaaaagt	cta aaaat	tcaat tg	tgttgcgc	actagagagt	ctccctgagg	5886
aagggctgga	aggttta	ct tcact	cacag ag	ttatttgt	tgaacactgt	aacatgctaa	5946
aatgtttaco	agaggga	tg cagca	cctaa ca	accctcac	aagtttaaaa	attcggggat	6006
gtccacaact	gatcaag	cgg tgtga	gaagg ga	ataggaga	agactggcac	: aaaatttctc	6066
acattcctaa	tgtgaat	ata tatat	ttaag tt	atttgcta	ttgtttcttt	gtttgtgagt	6126

		WARF023	35.ST25.txt		
ctttttggtt cctgccattg	tgattgcatg	taatttttt	ctagggttgt	ttgtttgttg	6186
agtetetete teattggatg	taattctctt	ttggtaacaa	attaacaatc	tatttgtatt	6246
atacgctttc agaatctatt	acttatttgt	aattgtttct	ttgtttgtaa	attgtgagta	6306
tcttattgta tggaattttc	tgattttatt	ttgaaaacaa	atcaataaga	tccatctgca	6366
ttatactccc ttcgtctcat	tttatgtgac	actttttgga	tttcgagatt	ctttgatctt	6426
aaatttttca tagatctttt	aaacattttg	agttatcaat	tattgtgatt	ttagtatttt	6486
ttatgtagtt tacaaataca	taaaatttat	ttttttaaa	aaaagaagat	ttcatgcgca	6546
aattcccgat caaacttaaa	ttactagact	ctcgaaaaat	gaaaagtgtc	acataaattg	6606
agacagaggg agtacttgtt	aatgttgtaa	ttattggcga	acaataatgt	tggtgattat	6666
cactttctga ataaatgttg	tgtcacgtgg	aaaaaacacc	aaatagaagt	attcatgctt	6726
ttttagtata tataaacatg	atttttaact	tggtttcagc	ggatagtcat	gacctttaac	6786
tctgaatgtg cacaagtaga	tacttgtata	aaattaaaca	aattttataa	aattatacaa	6846
tatgacactg agagtaattg	ataccaattg	cagtcgttgc	tgcttttcga	ttctctgtca	6906
ttctctaggt aattgatttt	acagaaaagg	gccaaaaata	tccctgaagt	accagaaaag	6966
gtctcaaaat accaaccatc	cacattttgg	tctaaaaata	tccttctact	catccttttt	7026
tgtctaaaat taccctttca	tccacatttt	tgctcactta	tacccttata	acaactctct	7086
ccttttttt aaaaaaatat	ttattatgtg	tcattttctt	attgaatgaa	ataaaaatcc	7146
acctctatta atttttccc	ataatttatc	caaatcaaaa	caatatattt	tttcaagatc	7206
caaaaaatat attttttaa	atctagtaat	ttctattttc	tatagctttt	ttccaaaaaa	7266
aaaatggttg ttttagataa	ttaaaatatc	tttaaaagta	ctagtcatgc	cacaattata	7326
gggacataat atattaatat	aaatcctaaa	atattttata	ataatatttt	attataaaat	7386
atattaatat attatgtccc	tgtaattgtg	gcatgactaa	tatttttaaa	aatattttaa	7446
ttatctaaaa caatttttt	tggaaaaaag	ctacagaaaa	tagaaattac	tagatttaaa	7506
aaaatacatt ttttggatct	tgaaaagata	tattgttttġ	atttggataa	attatgagaa	7566
aaaattaata gaggtggatt	tttatttcat	tcaataagaa	aatgacatat	aataagaatt	7626
taaaaaaaaa aaagagagag	ggttgttagt	tgcaagggta	taagtgtgca	aagtggtgga	7686
tggaagggta attttagacc	aaaaagatga	gtagaagggt	atttttagat	caaagatgga	7746
tggagggtat ttttagatct	tttcatgtac	ttcaggggta	tttttggccc	gattttattt	7806
gattctccct ctcttttgg	ttctggttga	ttgacaggcc	tgctaattga	gtgtttgttg	7866
taatcagtat taattactct	caaggtaata	ttatattcca	aacaaatttt	gtgttaccaa	7926
attaaatata tttctaaaac	tatcctgaag	gtagttaata	tactttttag	tgttgtatca	7986
tgtttttaat ataaaatatt	aaaatttaga	tgaaatttac	tttctagtta	aattggtcaa	8046
agttgtaaga atttcaagtg	aaagagtttt	taataatttc	acttttatgc	tatatatttt	8106
taaagttgaa cgactttta	ataaaaaaga	ataataaaat	tatatgataa	tttttataat	8166
acaatggcct ttatatgatg	aaaaaaaaga	aagaaattag	atgacaacaa	tgtccaaaaa	8226

WARF0235.ST25.txt	
taatcttaaa gaattatgat ttatatataa taaaattaaa tttaaaattt gatgaaaaaa	8286
tagagagaag aagaagatga tgaagtggaa ttatgtggtg gtgggtccat gtgacataat	8346
aaaaaacaat totottaaat aatootttoa tactaatgat aaaagaaaat atatatata	8406
atatatattt ctttttacaa attgtgaata gagaaaagga aaatggggta gcaattacaa	8466
ggtaggaaat caaactttat caacaagttg agagttcaag taatcaactt tatcatatcc	8526
gaaacattcc ttccgctttg agttcttttc tttatggatc ccg	8569
<210> 4 <211> 2913 <212> DNA <213> Solanum bulbocastanum	
<220> <221> CDS <222> (1)(2913)	
<pre>&lt;400&gt; 4 atg gct gaa gct ttc att caa gtt ctg cta gac aat ctc act tct ttc Met Ala Glu Ala Phe Ile Gln Val Leu Leu Asp Asn Leu Thr Ser Phe 1</pre>	48
ctc aaa ggg gaa ctt gta ttg ctt ttc ggt ttt caa gat gag ttc caa Leu Lys Gly Glu Leu Val Leu Leu Phe Gly Phe Gln Asp Glu Phe Gln 20 25 30	96
agg ctt tca agc atg ttt tct aca att caa gcc gtc ctt gaa gat gct Arg Leu Ser Ser Met Phe Ser Thr Ile Gln Ala Val Leu Glu Asp Ala 35 40 45	144
cag gag aag caa ctc aac aac cct cta gaa aat tgg ttg caa aaa Gln Glu Lys Gln Leu Asn Asn Lys Pro Leu Glu Asn Trp Leu Gln Lys 50 55 60	192
ctc aat gct gct aca tat gaa gtc gat gac atc ttg gat gaa tat aaa Leu Asn Ala Ala Thr Tyr Glu Val Asp Asp Ile Leu Asp Glu Tyr Lys 65 70 75 80	240
acc aag gcc aca aga ttc tcc cag tct gaa tat ggc cgt tat cat cca Thr Lys Ala Thr Arg Phe Ser Gln Ser Glu Tyr Gly Arg Tyr His Pro 85 90 95	288
aag gtt atc cct ttc cgt cac aag gtc ggg aaa agg atg gac caa gtg Lys Val Ile Pro Phe Arg His Lys Val Gly Lys Arg Met Asp Gln Val 100 105 110	336
atg aaa aaa cta aag gca att gct gag gaa aga aag aat ttt cat ttg Met Lys Lys Leu Lys Ala Ile Ala Glu Glu Arg Lys Asn Phe His Leu 115 120 125	384
cac gaa aaa att gta gag aga caa gct gtt aga cgg gaa aca ggt tct His Glu Lys Ile Val Glu Arg Gln Ala Val Arg Arg Glu Thr Gly Ser 130 135 140	432
gta tta acc gaa ccg cag gtt tat gga aga gac aaa gag aaa gat gag Val Leu Thr Glu Pro Gln Val Tyr Gly Arg Asp Lys Glu Lys Asp Glu 145 150 155 160	480
ata gtg aaa atc cta ata aac aat gtt agt gat gcc caa cac ctt tca Ile Val Lys Ile Leu Ile Asn Asn Val Ser Asp Ala Gln His Leu Ser 165 170 175	528
gtc ctc cca ata ctt ggt atg ggg gga tta gga aaa acg act ctt gcc Val Leu Pro Ile Leu Gly Met Gly Gly Leu Gly Lys Thr Thr Leu Ala	576

				gac Asp											624
				tcg Ser											672
-		-	-	att Ile 230	_								_	_	720
				aag Lys											768
				gat Asp											816
				gtc Val											864
				ctt Leu											912
				tca Ser 310											960
				ttt Phe											1008
				gag Glu											1056
_	-			gga Gly			_	-		_	_	-	-	_	1104
-		-		 aga Arg	-	_	-				_			_	1152
			_	cct Pro 390					_						1200
	-	_		tgc Cys				_					-	_	1248
-		_		gaa Glu	_						-				1296
				gga Gly		-				_			-	-	1344
_			_	tac Tyr	-								-	-	1392

	gat Asp								cat	gat		atc	cat			1440
_	aca Thr		_			_				_	_			_	-	1488
	aat Asn															1536
	gtg Val															1584
-	gtg Val 530					_	_				_					1632
	gga Gly															1680
_	cgt Arg	-			-	_		-	_				_			1728
	gat Asp				_		_		-	_	_			_		1776
_	aaa Lys			_		_					-		-	-		1824
_	act Thr 610	-	_							_		_		_		1872
	ggt Gly															1920
	gga Gly											-				1968
	gtg Val															2016
	aat Asn	Leu					Met									2064
		675					680					685				
	tat Tyr 690	gaa					aaa					ctc				2112
Ile	Tyr	gaa Glu ctg	Ser	Glu	Glu	Val 695 aaa	aaa Lys atc	Val	Leu ggc	Glu	Ala 700 aga	ctc Leu gga	Lys	Pro	His	2112
tcc Ser 705	Tyr 690 aat	gaa Glu ctg Leu	Ser act Thr	Glu tct Ser	Glu tta Leu 710 cac	Val 695 aaa Lys tca	aaa Lys atc Ile	Val tat Tyr	ggc Gly aaa	Glu ttc Phe 715 aat	Ala 700 aga Arg	ctc Leu gga Gly	Lys atc Ile tct	Pro cat His	His ctc Leu 720 cta	

			gag Glu					2304
			att Ile 775					2352
			agg Arg					2400
			aag Lys					2448
			gag Glu					2496
			ctc Leu					2544
			ttc Phe 855					2592
			aat Asn					2640
			agt Ser					2688
			Gly ggg					2736
			aac Asn					2784
			aca Thr 935					2832
			aag Lys					2880
			aat Asn		taa			2913

<210> 5 <211> 970 <212> PRT <213> Solanum bulbocastanum

<400> 5

Met Ala Glu Ala Phe Ile Gln Val Leu Leu Asp Asn Leu Thr Ser Phe 1 5 10 15

									WAR	F023	5.ST	'25.t	xt		
Leu	Lys	Gly	Glu 20	Leu	Val	Leu	Leu	Phe 25						Phe	Gln
Arg	Leu	Ser 35	Ser	Met	Phe	Ser	Thr 40	Ile	Gln	Ala	Val	Leu 45	Glu	Asp	Ala
Gln	Glu 50	Lys	Gln	Leu	Asn	Asn 55	Lys	Pro	Leu	Glu	Asn 60	Trp	Leu	Gln	Lys
Leu 65	Asn	Ala	Ala	Thr	Tyr 70	Glu	Val	Asp	Asp	Ile 75	Leu	Asp	Glu	Tyr	Lys 80
Thr	Lys	Ala	Thr	Arg 85	Phe	Ser	Gln	Ser	Glu 90	Tyr	Gly	Arg	Tyr	His 95	Pro
Lys	Val	Ile	Pro 100	Phe	Arg	His	Lys	Val 105	Gly	Lys	Arg	Met	Asp 110	Gln	Val
Met	Lys	Lys 115	Leu	Lys	Ala	Ile	Ala 120	Glu	Glu	Arg	Lys	Asn 125	Phe	His	Leu
His	Glu 130	Lys	Ile	Val	Glu	Arg 135	Gln	Ala	Val	Arg	Arg 140	Glu	Thr	Gly	Ser
Val 145	Leu	Thr	Glu	Pro	Gln 150	Val	Tyr	Gly	Arg	Asp 155	Lys	Glu	Lys	Asp	Glu 160
Ile	Val	Lys	Ile	Leu 165	Ile	Asn	Asn	Val	Ser 170	Asp	Ala	Gln	His	Leu 175	Ser
Val	Leu	Pro	Ile 180	Leu	Gly	Met	Gly	Gly 185	Leu	Gly	Lys	Thr	Thr 190	Leu	Ala
Gln	Met	Val 195	Phe	Asn	Asp	Gln	Arg 200	Val	Thr	Glu	His	Phe 205	His	Ser	Lys

Ile Trp Ile Cys Val Ser Glu Asp Phe Asp Glu Lys Arg Leu Ile Lys 210 215 220

Ala Ile Val Glu Ser Ile Glu Gly Arg Pro Leu Leu Gly Glu Met Asp 225 230 235 240

Leu Ala Pro Leu Gln Lys Lys Leu Gln Glu Leu Leu Asn Gly Lys Arg  $245 \hspace{1.5cm} 250 \hspace{1.5cm} 255$ 

Tyr Leu Leu Val Leu Asp Asp Val Trp Asn Glu Asp Gln Gln Lys Trp 260 265 270

Ala Asn Leu Arg Ala Val Leu Lys Val Gly Ala Ser Gly Ala Ser Val 275 280 285

Leu Thr Thr Thr Arg Leu Glu Lys Val Gly Ser Ile Met Gly Thr Leu 290  $\phantom{\bigg|}$  295  $\phantom{\bigg|}$  300  $\phantom{\bigg|}$ 

Gln 305	Pro	Tyr	Glu	Leu	Ser 310	Asn	Leu	Ser	Gln	Glu 315	Asp	Cys	Trp	Leu	Leu 320
Phe	Met	Gln	Arg	Ala 325	Phe	Gly	His	Gln	Glu 330	Glu	Ile	Asn	Pro	Asn 335	Leu
Val	Ala	Ile	Gly 340	Lys	Glu	Ile	Val	Lys 345	Lys	Ser	Gly	Gly	Val 350	Pro	Leu
Ala	Ala	Lys 355	Thr	Leu	Gly	Gly	Ile 360	Leu	Cys	Phe	Lys	Arg 365	Glu	Glu	Arg
Ala	Trp 370	Glu	His	Val	Arg	Asp 375	Ser	Pro	Ile	Trp	Asn 380	Leu	Pro	Gln	Asp
Glu 385	Ser	Ser	Ile	Leu	Pro 390	Ala	Leu	Arg	Leu	Ser 395	Tyr	His	Gln	Leu	Pro 400
Leu	Asp	Leu	Lys	Gln 405	Cys	Phe	Ala	Tyr	Cys 410	Ala	Val	Phe	Pro	Lys 415	Asp
Ala	Lys	Met	Lys 420	Lys	Glu	Lys	Leu	Ile 425	Ser	Leu	Trp	Met	Ala 430	His	Gly
Phe	Leu	Leu 435	Ser	Lys	Gly	Asn	Met 440	Glu	Leu	Glu	Asp	Val 445	Gly	Asp	Glu
Val	Trp 450	Lys	Glu	Leu	Tyr	Leu 455	Arg	Ser	Phe	Phe	Gln 460	Glu	Ile	Glu	Val
Lys 465	Asp	Gly	Lys	Thr	Туг 470	Phe	Lys	Met	His	Asp 475	Leu	Ile	His	Asp	Leu 480
Ala	Thr	Ser	Leu	Phe 485	Ser	Ala	Asn	Thr	Ser 490	Ser	Ser	Asn	Ile	Arg 495	Glu
Ile	Asn	Lys	His 500		Туг	Thr	His	Met 505		Ser	Ile	Gly	Phe 510		Glu
Val	Val	Phe 515	Phe	Tyr	Thr	Leu	Pro 520	Pro	Leu	Glu	Lys	Phe 525	Ile	Ser	Leu
Arg	Val 530	Leu	Asn	Leu	Gly	Asp 535	Ser	Thr	Phe	Asn	Lys 540	Leu	Pro	Ser	Ser
Ile 545	Gly	Asp	Leu	Val	His 550	Leu	Arg	Tyr	Leu	Asn 555	Leu	Tyr	Gly	Ser	Gly 560
Met	Arg	Ser	Leu	Pro	Lys	Gln	Leu	Cys	Lys 570	Leu	Gln	Asn	Leu	Gln 575	Thr

Leu	Asp	Leu	Gln 580	Tyr	Cys	Thr	Lys	Leu 585			5.ST Leu		-	Glu	Thr
Ser	Lys	Leu 595	Gly	Ser	Leu	Arg	Asn 600	Leu	Leu	Leu	Asp	Gly 605	Ser	Gln	Ser
Leu	Thr 610	Cys	Met	Pro	Pro	Arg 615	Ile	Gly	Ser	Leu	Thr 620	Cys	Leu	Lys	Thr
Leu 625	Gly	Gln	Phe	Val	Val 630	Gly	Arg	Lys	Lys	Gly 635	Tyr	Gln	Leu	Gly	Glu 640
Leu	Gly	Asn	Leu	Asn 645	Leu	Tyr	Gly	Ser	Ile 650	Lys	Ile	Ser	His	Leu 655	Glu
Arg	Val	Lys	Asn 660	Asp	Met	Asp	Ala	Lys 665	Glu	Ala	Asn	Leu	Ser 670	Ala	Lys
Gly	Asn	Leu 675	His	Ser	Leu	Ser	Met 680	Ser	Trp	Asn	Asn	Phe 685	Gly	Pro	His
Ile	Tyr 690	Glu	Ser	Glu	Glu	Val 695	Lys	Val	Leu	Glu	Ala 700	Leu	Lys	Pro	His
Ser 705	Asn	Leu	Thr	Ser	Leu 710	Lys	Ile	Tyr	Gly	Phe 715	Arg	Gly	Ile	His	Leu 720
Pro	Glu	Trp	Met	Asn 725	His	Ser	Val	Leu	Lys 730	Asn	Ile	Val	Ser	Ile 735	Leu
Ile	Ser	Asn	Phe 740	Arg	Asn	Cys	Ser	Cys 745	Leu	Pro	Pro	Phe	Gly 750	Asp	Leu
Pro	Cys	Leu 755	Glu	Ser	Leu	Glu	Leu 760	His	Trp	Gly	Ser	Ala 765	Asp	Val	Glu
Tyr	Val 770	Glu	Glu	Val	Asp	Ile 775	Asp	Val	His	Ser	Gly 780	Phe	Pro	Thr	Arg
Ile 785	Arg	Phe	Pro	Ser	Leu 790	Arg	Lys	Leu	Asp	Ile 795	Trp	Asp	Phe	Gly	Ser 800
Leu	Lys	Gly	Leu	Leu 805	Lys	Lys	Glu	Gly	Glu 810	Glu	Gln	Phe	Pro	Val 815	Leu
Glu	Glu	Met	Ile 820	Ile	His	Glu	Cys	Pro 825	Phe	Leu	Thr	Leu	Ser 830	Ser	Asn
Leu	Arg	Ala 835	Leu	Thr	Ser	Leu	Arg 840	Ile	Cys	Tyr	Asn	Lys 845	Val	Ala	Thr
_															

Ser Phe Pro Glu Glu Met Phe Lys Asn Leu Ala Asn Leu Lys Tyr Leu 850 860

Thr 865	Ile	Ser	Arg	Cys	Asn 870	Asn	Leu	Lys	Glu	Leu 875	Pro	Thr	Ser	Leu	Ala 880	
Ser	Leu	Asn	Ala	Leu 885	Lys	Ser	Leu	Lys	Ile 890	Gln	Leu	Cys	Cys	Ala 895	Leu	
Glu	Ser	Leu	Pro 900	Glu	Glu	Gly	Leu	Glu 905	Gly	Leu	Ser	Ser	Leu 910	Thr	Glu	
Leu	Phe	Val 915	Glu	His	Cys	Asn	Met 920	Leu	Lys	Cys	Leu	Pro 925	Glu	Gly	Leu	
Gln	His 930	Leu	Thr	Thr	Leu	Thr 935	Ser	Leu	Lys	Ile	Arg 940	Gly	Cys	Pro	Gln	
Leu 945	Ile	Lys	Arg	Cys	Glu 950	Lys	Gly	Ile	Gly	Glu 955	Asp	Trp	His	Lys	Ile 960	
Ser	His	Ile	Pro	Asn 965	Val	Asn	Ile	Tyr	Ile 970							
<210 <211 <212 <213	.> 3 ?> [	5 3592 DNA Solar	num k	oulbo	ocast	canur	n									
<220 <221 <222	.> e	exon	. (413	L)							٠					
<220 <221 <222	.> ]	Intro (412)	on	549)												
<220 <221 <222	.> €	exon (650)	(3	3592)	ı											
	gct	gaa			att Ile	Gln	Val	Leu		Āsp	Asn	Leu	Thr	Ser		48
					gta Val											96
					ttt Phe											144
_		_			aac Asn		_			_			_			192
		-	_		tat Tyr 70	-	-	-	_		_	-	-			240
acc	aag	gcc	aca	aga	ttc	tcc	cag	tct	gaa	tat	ggc	cgt	tat	cat	cca	288

WARF0235.ST25.txt Thr Lys Ala Thr Arg Phe Ser Gln Ser Glu Tyr Gly Arg Tyr His Pro 85 90 95	
aag gtt atc cct ttc cgt cac aag gtc ggg aaa agg atg gac caa gtg Lys Val Ile Pro Phe Arg His Lys Val Gly Lys Arg Met Asp Gln Val 100 105 110	336
atg aaa aaa cta aag gca att gct gag gaa aga aag aat ttt cat ttg Met Lys Lys Leu Lys Ala Ile Ala Glu Glu Arg Lys Asn Phe His Leu 115 120 125	384
cac gaa aaa att gta gag aga caa gct gttagacggg aaacaggtac His Glu Lys Ile Val Glu Arg Gln Ala 130 135	431
tcatcttaaa ttagtattac aacaactaag tttatattca tttttttggc aattatgaaa	491
ttcagaaaag ggttaaatat actcatgtcc tatcgtaaat agtgtaaata tacctctcgt	551
tgtactttcg atctgaatat acttgtcaaa tctggcaagc tcagaatcaa attatccacc	611
ccaactttta aatactcgat atctttagaa atccacct gtc taa ctc atc cac tac Val Leu Ile His Tyr 140	667
cca ttc cct ttg ctt tga att ctt ttc ttt acc tat aaa cgt gga aca Pro Phe Pro Leu Leu Ile Leu Phe Phe Thr Tyr Lys Arg Gly Thr 145 150 155	715
ctc gat ccg ttt tgc ttt tct taa caa agc agc tca gag aaa aga ggt Leu Asp Pro Phe Cys Phe Ser Gln Ser Ser Ser Glu Lys Arg Gly 160 165 170	763
ttt ctt cta ttc tgt ttc tct gtg tgc tgc act tgg gtc ctt aat ccc Phe Leu Phe Cys Phe Ser Val Cys Cys Thr Trp Val Leu Asn Pro 175 180 185	811
att aaa aac agg gca tgt taa tcc caa cga cgg tag cct ttc ctg aca Ile Lys Asn Arg Ala Cys Ser Gln Arg Arg Pro Phe Leu Thr 190 195 200	859
gct gac tgt aaa ttt tgt cta aca aag aaa aaa aaa gat tag aca tgt Ala Asp Cys Lys Phe Cys Leu Thr Lys Lys Lys Asp Thr Cys 205 210 215	907
ttt tcc ttg tca ttg att agg ctg gat ttc ttt cag agt gga aca tag Phe Ser Leu Ser Leu Ile Arg Leu Asp Phe Phe Gln Ser Gly Thr 220 225 230	955
ggg ata tat tgg acc aaa agt aga atg ggt ata tat tta aag tat ttc Gly Ile Tyr Trp Thr Lys Ser Arg Met Gly Ile Tyr Leu Lys Tyr Phe 235 240 245	1003
tga tag aac agg agt ata ttg tgc gaa aat atc ctc tat ttt ctg ttg Asn Arg Ser Ile Leu Cys Glu Asn Ile Leu Tyr Phe Leu Leu 250 . 255 . 260	1051
tct cct aat gag ttt gaa tgt aat aat att ctc atg tgg aca ttg ctt Ser Pro Asn Glu Phe Glu Cys Asn Asn Ile Leu Met Trp Thr Leu Leu 265 270 275	1099
gca cca ggt tct gta tta acc gaa ccg cag gtt tat gga aga gac aaa Ala Pro Gly Ser Val Leu Thr Glu Pro Gln Val Tyr Gly Arg Asp Lys 280 285 290	1147
gag aaa gat gag ata gtg aaa atc cta ata aac aat gtt agt gat gcc Glu Lys Asp Glu Ile Val Lys Ile Leu Ile Asn Asn Val Ser Asp Ala 295 300 305 310	1195
caa cac ctt tca gtc ctc cca ata ctt ggt atg ggg gga tta gga aaa	1243

Gln	His	Leu	Ser	Val 315	Leu	Pro	Ile	Leu			Gly			Gly 325	Lys	
								aat Asn 335								1291
								gtc Val								1339
			_	-		_	_	tct Ser		_						1387
								caa Gln								1435
aat Asn	gga Gly	aaa Lys	aga Arg	tac Tyr 395	ttg Leu	ctt Leu	gtc Val	tta Leu	gat Asp 400	gat Asp	gtt Val	tgg Trp	aat Asn	gaa Glu 405	gat Asp	1483
								gca Ala 415								1531
ggt Gly	gct Ala	tct Ser 425	gtt Val	cta Leu	acc Thr	act Thr	act Thr 430	cgt Arg	ctt Leu	gaa Glu	aag Lys	gtt Val 435	gga Gly	tca Ser	att Ile	1579
								ctg Leu								1627
								gca Ala								1675
aat Asn	cca Pro	aac Asn	ctt Leu	gtg Val 475	gca Ala	atc Ile	gga Gly	aag Lys	gag Glu 480	att Ile	gtg Val	aaa Lys	aaa Lys	agt Ser 485	ggt Gly	1723
								ctt Leu 495								1771
						Ğlu		gtg Val								1819
								ctg Leu								1867
								caa Gln								1915
								aaa Lys								1963
-								aaa Lys 575			-	-			_	2011
								tta Leu								2059

		gaa Glu														2107
		gat Asp	_	-			_			-				_		2155
		cgt Arg	-					_				_	_			2203
		gcc Ala														2251
		tcg Ser 665		_						_	_				-	2299
		tct Ser				_		-			_		_		_	2347
		agt Ser														2395
	_	caa Gln			_				-		_		_	_	_	2443
		gaa Glu		_				_		_					_	2491
		cag Gln 745														2539
_		aag Lys						-	-			_				2587
		ggt Gly	-													2635
		ctt Leu														2683
		gca Ala														2731
		cca Pro 825				_		_	_	-				_	_	2779
		cca Pro														2827
		cat His														2875
gtc	tct	att	cta	att	agc	aac	ttc	aga	aac	tgc	tca	tgc	tta	сса	ссс	2923

Val Ser	Ile Leu	ı Ile Sei 875	Asn	Phe	_			Ser C		Pro 885		
ttt ggt Phe Gly		Pro Cys		Glu						Gĺy		2971
gcg gat Ala Asp								Asp V				3019
ttc ccc Phe Pro 920	aca aga Thr Arc	ata ago Ile Aro	ttt Phe 925	cca Pro	tcc Ser	ttg Leu	Arg :	aaa c Lys Le 930	tt gat eu Asp	ata Ile	tgg Trp	3067
gac ttt Asp Phe 935			Gly			Lys						3115
ttc cct Phe Pro					Ile :						Thr	3163
ctt tct Leu Ser		Leu Arg								Tyr		3211
aaa gta Lys Val								Lys A:				3259
ctc aaa Leu Lys 1000	Tyr Le	g aca at eu Thr Il		r Ar					Lys		_	3304
cct acc Pro Thr 1015	Ser Le	g gct ag u Ala Se		ı As	_		-	a agt s Ser 102!	Leu			3349
caa ttg Gln Leu 1030	Cys Cy	rc gca ct vs Ala Le		ı Se			-	g gaa u Glu 1040	Gly	_		3394
ggt tta Gly Leu 1045	Ser Se	a ctc ac r Leu Th		ı Le				a cac u His 105!	Cys		_	3439
cta aaa Leu Lys 1060	Cys Le	a cca ga u Pro Gl		y Le				a aca u Thr 1070	acc Thr			3484
agt tta Ser Leu 1075	Lys Il	t cgg gg e Arg Gl		s Pr					Arg			3529
aag gga Lys Gly 1090	Ile ĞÎ	a gaa ga y Glu As		Hi					Ile			3574
gtg aat Val Asn 1105	Ile Ty		ıa									3592
<212> D	913 NA	bulbocas	tanur	n								

.000									WAR	F023	5.ST	'25.t	xt			
<220 <221 <222	.> (	CDS (1).	. (29)	13)												
	gct	gaa						ctg Leu								48
								ttc Phe 25								96
								att Ile								144
								cct Pro								192
								gat Asp								240
								tct Ser								288
								gtc Val 105								336
								gag Glu								384
								gct Ala								432
								gga Gly								480
								gtt Val	-	_	-					528
								gga Gly 185								576
	-	-			-	_	_	gtt Val								624
			_	-	_	-	-	ttt Phe	_		-				_	672
-		_	_			_		agg Arg						_	_	720
								cag Gln								768

Page 27

tac ttg ctt gtc tta gat gat gtt tgg aat gaa gat caa cag aag tgg 816

Tyr	Leu	Leu	Val 260	Leu	Asp	Asp	Val	Trp 265			Asp			Lys	Trp	
-			_	-	gtc Val	_	_	-		-	_	-	_		_	864
					ctt Leu											912
			-	_	tca Ser 310		_			_	_	_			_	960
					ttt Phe											1008
					gag Glu											1056
gca Ala	gcc Ala	aaa Lys 355	act Thr	ctt Leu	gga Gly	ggt Gly	att Ile 360	ttg Leu	tgc Cys	ttc Phe	aag Lys	aga Arg 365	gaa Glu	gaa Glu	aga Arg	1104
					aga Arg											1152
					cct Pro 390											1200
					tgc Cys											1248
		_	_		gaa Glu	_						_				1296
					gga Gly											1344
					tac Tyr											1392
					tat Tyr 470											1440
					tca Ser											1488
					tac Tyr											1536
					act Thr											1584
					ggt Gly	_	-				_					1632

					cat His 550											1680
					aag Lys											1728
					tgc Cys											1776
					ctc Leu											1824
_		_			cca Pro					_		-				1872
				-	gtt Val 630			_							_	1920
					ctc Leu											1968
					aag Lys											2016
					tta Leu											2064
		-		-	gaa Glu	-				_	_					2112
					tta Leu 710											2160
					cac His											2208
att Ile	agc Ser	aac Asn	ttc Phe 740	aga Arg	aac Asn	tgc Cys	tca Ser	tgc Cys 745	tta Leu	cca Pro	ccc Pro	ttt Phe	ggt Gly 750	gat Asp	ctg Leu	2256
					cta Leu											2304
	-	-	-		gat Asp		-	-							_	2352
					ttg Leu 790											2400
					aaa Lys											2448
gaa	gag	atg	ata	att	cac	gag	tgc	cct	ttt	ctg	acc	ctt	tct	tct	aat	2496

Glu Glu Met Ile Ile His Glu Cys Pro Phe Leu Thr Leu Ser Ser Asn 820 825 830	
ctt agg gct ctt act tcc ctc aga att tgc tat aat aaa gta gct act Leu Arg Ala Leu Thr Ser Leu Arg Ile Cys Tyr Asn Lys Val Ala Thr 835 840 845	2544
tca ttc cca gaa gag atg ttc aaa aac ctt gca aat ctc aaa tac ttg Ser Phe Pro Glu Glu Met Phe Lys Asn Leu Ala Asn Leu Lys Tyr Leu 850 855 860	2592
aca atc tct cgg tgc aat aat ctc aaa gag ctg cct acc agc ttg gct Thr Ile Ser Arg Cys Asn Asn Leu Lys Glu Leu Pro Thr Ser Leu Ala 865 870 875 880	2640
agt ctg aat gct ttg aaa agt cta aaa att caa ttg tgt tgc gca cta Ser Leu Asn Ala Leu Lys Ser Leu Lys Ile Gln Leu Cys Cys Ala Leu 885 890 895	2688
gag agt ctc cct gag gaa ggg ctg gaa ggt tta tct tca ctc aca gag Glu Ser Leu Pro Glu Glu Gly Leu Glu Gly Leu Ser Ser Leu Thr Glu 900 905 910	2736
tta ttt gtt gaa cac tgt aac atg cta aaa tgt tta cca gag gga ttg Leu Phe Val Glu His Cys Asn Met Leu Lys Cys Leu Pro Glu Gly Leu 915 920 925	2784
cag cac cta aca acc ctc aca agt tta aaa att cgg gga tgt cca caa Gln His Leu Thr Thr Leu Thr Ser Leu Lys Ile Arg Gly Cys Pro Gln 930 935 940	2832
ctg atc aag cgg tgt gag aag gga ata gga gaa gac tgg cac aaa att Leu Ile Lys Arg Cys Glu Lys Gly Ile Gly Glu Asp Trp His Lys Ile 945 950 955 960	2880
tct cac att cct aat gtg aat ata tat att taa Ser His Ile Pro Asn Val Asn Ile Tyr Ile 965 970	2913
<210> 8 <211> 970 <212> PRT <213> Solanum bulbocastanum	
<400> 8	
Met Ala Glu Ala Phe Ile Gln Val Leu Leu Asp Asn Leu Thr Ser Phe 1 5 10 15	
Leu Lys Gly Glu Leu Val Leu Leu Phe Gly Phe Gln Asp Glu Phe Gln 20 25 30	
Arg Lou Cor Cor Mot Dho Cor Mbr Tle Cla Ale Wel Lou Clu Arg Ale	
Arg Leu Ser Ser Met Phe Ser Thr Ile Gln Ala Val Leu Glu Asp Ala 35 40 45	
•	
35 40 45  Gln Glu Lys Gln Leu Asn Asn Lys Pro Leu Glu Asn Trp Leu Gln Lys	

•	Lys	Val	Ile	Pro 100	Phe	Arg	His	Lys	Val 105	Gly	Lys	Arg	Met	Asp 110	Gln	Val
	Met	Lys	Lys 115	Leu	Lys	Ala	Ile	Ala 120	Glu	Glu	Arg	Lys	Asn 125	Phe	His	Leu
	His	Glu 130	Lys	Ile	Val	Glu	Arg 135	Gln	Ala	Val	Arg	Arg 140	Glu	Thr	Gly	Ser
	Val 145	Leu	Thr	Glu	Pro	Gln 150	Val	Tyr	Gly	Arg	Asp 155	Lys	Glu	Lys	Asp	Glu 160
	Ile	Val	Lys	Ile	Leu 165	Ile	Asn	Asn	Val	Ser 170	Asp	Ala	Gln	His	Leu 175	Ser
	Val	Leu	Pro	Ile 180	Leu	Gly	Met	Gly	Gly 185	Leu	Gly	Lys	Thr	Thr 190	Leu	Ala
	Gln	Met	Val 195	Phe	Asn	Asp	Gln	Arg 200	Val	Thr	Glu	His	Phe 205	His	Ser	Lys
	Ile	Trp 210	Ile	Суѕ	Val	Ser	Glu 215	Asp	Phe	Asp	Glu	Lys 220	Arg	Leu	Ile	Lys
	Ala 225	Ile	Val	Glu	Ser	Ile 230	Glu	Gly	Arg	Pro	Leu 235	Leu	Gly	Glu	Met	Asp 240
	Leu	Ala	Pro	Leu	Gln 245	Lys	Lys	Leu	Gln	Glu 250	Leu	Leu	Asn	Gly	Lys 255	Arg
	Tyr	Leu	Leu	Val 260	Leu	Asp	Asp	Val	Trp 265	Asn	Glu	Asp	Gln	Gln 270	Lys	Trp
	Ala	Asn	Leu 275	Arg	Ala	Val	Leu	Lys 280	Val	Gly	Ala	Ser	Gly 285	Ala	Ser	Val
	Leu	Thr 290	Thr	Thr	Arg	Leu	Glu 295	Lys	Val	Gly	Ser	Ile 300	Met	Gly	Thr	Leu
	Gln 305	Pro	Tyr	Glu	Leu	Ser 310	Asn	Leu	Ser	Gln	Glu 315	Asp	Cys	Trp	Leu	Leu 320
	Phe	Met	Gln	Arg	Ala 325	Phe	Gly	His	Gln	Glu 330	Glu	Ile	Asn	Pro	Asn 335	Leu
	Val	Ala	Ile	Gly 340	Lys	Glu	Ile	Val	Lys 345	Lys	Ser	Gly	Gly	Val 350	Pro	Leu
	Ala	Ala	Lys 355	Thr	Leu	Gly	Gly	Ile 360	Leu	Cys	Phe	Lys	Arg 365	Glu	Glu	Arg
	Ala	Trp	Glu	His	Val	Arg	Asp	Ser	Pro	Ile	_	Asn ge 3	_	Pro	Gln	Asp

375

Glu 385	Ser	Ser	Ile	Leu	Pro 390	Ala	Leu	Arg	Leu	Ser 395	Tyr	His	Gln	Leu	Pro 400
Leu	Asp	Leu	Lys	Gln 405	Cys	Phe	Ala	Tyr	Cys 410	Ala	Val	Phe	Pro	Lys 415	Asp
Ala	Lys	Met	Glu 420	Lys	Glu	Lys	Leu	Ile 425	Ser	Leu	Trp	Met	Ala 430	His	Gly
Phe	Leu	Leu 435	Ser	Lys	Gly	Asn	Met 440	Glu	Leu	Glu	Asp	Val 445	Gly	Asp	Glu
Val	Trp 450	Lys	Glu	Leu	Tyr	Leu 455	Arg	Ser	Phe	Phe	Gln 460	Glu	Ile	Glu	Val
Lys 465	Asp	Gly	Lys	Thr	Tyr 470	Phe	Lys	Met	His	Asp 475	Leu	Ile	His	Asp	Leu 480
Ala	Thr	Ser	Leu	Phe 485	Ser	Ala	Asn	Thr	Ser 490	Ser	Ser	Asn	Ile	Arg 495	Glu
Ile	Asn	Lys	His 500	Ser	Tyr	Thr	His	Met 505	Met	Ser	Ile	Gly	Phe 510	Ala	Glu
Val	Val	Phe 515	Phe	Tyr	Thr	Leu	Pro 520	Pro	Leu	Glu	Lys	Phe 525	Ile	Ser	Leu
Arg	Val 530	Leu	Asn	Leu	Gly	Asp 535	Ser	Thr	Phe	Asn	Lys 540	Leu	Pro	Ser	Ser
Ile 545	Gly	Asp	Leu	Val	His 550	Leu	Arg	Tyr	Leu	Asn 555	Leu	Tyr	Gly	Ser	Gly 560
Met	Arg	Ser	Leu	Pro 565	Lys	Gln	Leu	Cys	Lys 570	Leu	Gln	Asn	Leu	Gln 575	Thr
Leu	Asp	Leu	Gln 580	Tyr	Cys	Thr	Lys	Leu 585	Cys	Cys	Leu	Pro	Lys 590	Glu	Thr
Ser	Lys	Leu 595	Gly	Ser	Leu	Arg	Asn 600	Leu	Leu	Leu	Asp	Gly 605	Ser	Gln	Ser
Leu	Thr 610	Cys	Met	Pro	Pro	Arg 615	Ile	Gly	Ser	Leu	Thr 620	Cys	Leu	Lys	Thr
Leu 625	Gly	Gln	Phe	Val	Val 630	Gly	Arg	Lys	Lys	Gly 635	Tyr	Gln	Leü	Gly	Glu 640
Leu	Gly	Asn	Leu	Asn 645	Leu	Tyr	Gly	Ser	Ile 650	Lys	Ile	Ser	His	Leu 655	Glu

Arg	Val	Lys	Asn 660	Asp	Lys	Asp	Ala	Lys 665	Glu	Ala	Asn	Leu	Ser 670	Ala	Lys
Gly	Asn	Leu 675	His	Ser	Leu	Ser	Met 680	Ser	Trp	Asn	Asn	Phe 685	Gly	Pro	His
Ile	Tyr 690	Glu	Ser	Glu	Glu	Val 695	Lys	Val	Leu	Glu	Ala 700	Leu	Lys	Pro	His
Ser 705	Asn	Leu	Thr	Ser	Leu 710	Lys	Ile	Tyr	Gly	Phe 715	Arg	Gly	Ile	His	Leu 720
Pro	Glu	Trp	Met	Asn 725	His	Ser	Val	Leu	Lys 730	Asn	Ile	Val	Ser	Ile 735	Leu
Ile	Ser	Asn	Phe 740	Arg	Asn	Cys	Ser	Cys 745	Leu	Pro	Pro	Phe	Gly 750	Asp	Leu
Pro	Cys	Leu 755	Glu	Ser	Leu	Glu	Leu 760	His	Trp	Gly	Ser	Ala 765	Asp	Val	Glu
Tyr	Val 770	Glu	Glu	Val	Asp	Ile 775	Asp	Val	His	Ser	Gly 780	Phe	Pro	Thr	Arg
Ile 785	Arg	Phe	Pro	Ser	Leu 790	Arg	Lys	Leu	Asp	Ile 795	Trp	Asp	Phe	Gly	Ser 800
Leu	Lys	Gly	Leu	Leu 805	Lys	Lys	Glu	Gly	Glu 810	Glu	Gln	Phe	Pro	Val 815	Leu
Glu	Glu	Met	Ile 820	Ile	His	Glu	Cys	Pro 825	Phe	Leu	Thr	Leu	Ser 830	Ser	Asn
Leu	Arg	Ala 835	Leu	Thr	Ser	Leu	Arg 840	Ile	Cys	Tyr	Asn	Lys 845	Val	Ala	Thr
Ser	Phe 850	Pro	Glu	Glu	Met	Phe 855	Lys	Asn	Leu	Ala	Asn 860	Leu	Lys	Tyr	Leu
Thr 865	Ile	Ser	Arg	Cys	Asn 870	Asn	Leu	Lys	Glu	Leu 875	Pro	Thr	Ser	Leu	Ala 880
Ser	Leu	Asn	Ala	Leu 885	Lys	Ser	Leu	Lys	Ile 890	Gln	Leu	Cys	Суз	Ala 895	Leu
Glu	Ser	Leu	Pro 900	Glu	Glu	Gly	Leu	Glu 905	Gly	Leu	Ser	Ser	Leu 910	Thr	Glu
Leu	Phe	Val 915	Glu	His	Cys	Asn	Met 920	Leu	Lys	Cys	Leu	Pro 925	Glu	Gly	Leu
Gln	His	Leu	Thr	Thr	Leu	Thr	Ser	Leu	Lys		Arg ge 3		Cys	Pro	Gln

930 935 940

Leu Ile Lys	Arg Cys Glu Lys	Gly Ile Gly Glu Asp T	frp His Lys Ile
945	950	955	960

Ser His Ile Pro Asn Val Asn Ile Tyr Ile 965 970

<210 <211 <212 <213	11> 2979 12> DNA															
<220 <220 <220	L>	CDS (1).	. (297	79)												
	gct	9 gaa Glu														48
		ggg Gly														96
		tca Ser 35														144
		aag Lys														192
		gtt Val														240
		gca Ala														288
		atc Ile														336
		aaa Lys 115														384
		agg Arg														432
		act Thr	Glu		Lys	Val			Arg		Lys					480
		aaa Lys														528
		cca Pro														576

					gat Asp				act		cat	ttc	aat			624
					tca Ser											672
					att Ile 230											720
					aag Lys											768
					gat Asp											816
aat Asn	ctt Leu	aga Arg 275	gca Ala	gta Val	ttg Leu	aag Lys	att Ile 280	gga Gly	gct Ala	agt Ser	ggt Gly	gct Ala 285	tca Ser	att Ile	cta Leu	864
					gaa Glu											912
		-			aat Asn 310	_			_	-	_		_	_		960
_		_	_		tgc Cys				_		_				_	1008
					att Ile											1056
					ggc Gly											1104
					gat Asp											1152
aat Asn 385	tct Ser	gtt Val	ttg Leu	cct Pro	gcc Ala 390	ctg Leu	agg Arg	ctg Leu	agt Ser	tat Tyr 395	cat His	cat His	ctt Leu	cca Pro	ctt Leu 400	1200
					ttt Phe											1248
					tat Tyr											1296
					aac Asn	_				-				-	_	1344
					ttg Leu											1392
					ttc Phe											1440

480

465					4/0					4/5					480	
aca Thr	tct Ser	atg Met	ttt Phe	tca Ser 485	gca Ala	agc Ser	gca Ala	tca Ser	agc Ser 490	aga Arg	agt Ser	ata Ile	cgc Arg	caa Gln 495	ata Ile	1488
					gaa Glu											1536
gat Asp	atg Met	atg Met 515	tcc Ser	att Ile	ggt Gly	ttc Phe	tcc Ser 520	gaa Glu	gtg Val	gtg Val	tct Ser	tct Ser 525	tac Tyr	tct Ser	cct Pro	1584
					ttt Phe											1632
	-		-	_	tta Leu 550	_			-		-		_			1680
					tct Ser											1728
					aat Asn											1776
					ccg Pro											1824
					cac His											1872
	_	_		-	ctt Leu 630	_						_	_			1920
					ctt Leu		_		_					_		1968
					cat His											2016
					tct Ser											2064
_		_	_		aac Asn	_		-		-	-	_				2112
_	-				cat His 710			_				-			_	2160
	-			-	ctc Leu		_		_				_	_		2208
	-	-			cta Leu		-		-	-		_	_	_		2256

									WAR	F023	5.ST	'25.t	xt			
			ggt Gly													2304
			gag Glu													2352
			cca Pro													2400
ctg Leu	aaa Lys	gga Gly	ttg Leu	cag Gln 805	aga Arg	atg Met	aaa Lys	gga Gly	gca Ala 810	gag Glu	caa Gln	ttc Phe	ccc Pro	gtg Val 815	ctt Leu	2448
-		_	aag Lys 820		_	_	_		_		_		_			2496
			aag Lys													2544
			ata Ile													2592
			aca Thr													2640
			ata Ile													2688
			agc Ser 900													2736
			tac Tyr													2784
			ctc Leu													2832
-			gag Glu		-	-							-			2880
			tgt Cys													2928
			cac His 980													2976
taa									-							2979

<210> 10 <211> 992 <212> PRT <213> Solanum bulbocastanum

<400> 10

Met 1	Ala	Glu	Ala	Phe 5	Leu	Gln	Val	Leu			Asn			Phe 15	Phe
Ile	Gln	Gly	Glu 20	Leu	Gly	Leu	Val	Phe 25	Gly	Phe	Glu	Lys	Glu 30	Phe	Lys
Lys	Leu	Ser 35	Ser	Met	Phe	Ser	Met 40	Ile	Gln	Ala	Val	Leu 45	Glu	Asp	Ala
Gln	Glu 50	Lys	Gln	Leu	Lys	Tyr 55	Lys	Ala	Ile	Lys	Asn 60	Trp	Leu	Gln	Lys
Leu 65	Asn	Val	Ala	Ala	Tyr 70	Glu	Val	Asp	Asp	Ile 75	Leu	Asp	Asp	Cys	Lys 80
Thr	Glu	Ala	Ala	Arg 85	Phe	Lys	Gln	Ala	Val 90	Leu	Gly	Arg	Tyr	His 95	Pro
Arg	Thr	Ile	Thr 100	Phe	Cys	Tyr	Lys	Val 105	Gly	Lys	Arg	Met	Lys 110	Glu	Met
Met	Glu	Lys 115	Leu	Asp	Ala	Ile	Ala 120	Glu	Glu	Arg	Arg	Asn 125	Phe	His	Leu
Asp	Glu 130	Arg	Ile	Ile	Glu	Arg 135	Gln	Ala	Ala	Arg	Arg 140	Gln	Thr	Gly	Phe
Val 145	Leu	Thr	Glu	Pro	Lys 150	Val	Tyr	Gly	Arg	Glu 155	Lys	Glu	Glu	Asp	Glu 160
Ile	Val	Lys	Ile	Leu 165	Ile	Asn	Asn	Val	Ser 170	Tyr	Ser	Glu	Glu	Val 175	Pro
Val	Leu	Pro	Ile 180	Leu	Gly	Met	Gly	Gly 185	Leu	Gly	Lys	Thr	Thr 190	Leu	Ala
Gln	Met	Val 195	Phe	Asn	Asp		Arg 200		Thr	Glu	His	Phe 205	Asn	Leu	Lys
Ile	Trp 210	Val	Cys	Val	Ser	Asp 215	Asp	Phe	Asp	Glu	Lys 220	Arg	Leu	Ile	Lys
Ala 225	Ile	Val	Glu	Ser	Ile 230	Glu	Gly	Lys	Ser	Leu 235	Gly	Asp	Met	Asp	Leu 240
Ala	Pro	Leu	Gln	Lys 245	Lys	Leu	Gln	Glu	Leu 250	Leu	Asn	Gly	Lys	Arg 255	Tyr
Phe	Leu	Val	Leu 260	Asp	Asp	Val	Trp	Asn 265	Glu	Asp	Gln	Glu	Lys 270	Trp	Asp
Asn	Leu	Arg 275	Ala	Val	Leu	Lys	Ile 280	Gly	Ala	Ser	Gly	Ala 285	Ser	Ile	Leu

Ile	Thr 290	Thr	Arg	Leu	Glu	Lys 295	Ile	Gly	Ser	Ile	Met 300	Gly	Thr	Leu	Gln
Leu 305	Tyr	Gln	Leu	Ser	Asn 310	Leu	Ser	Gln	Glu	Asp 315	Cys	Trp	Leu	Leu	Phe 320
Lys	Gln	Arg	Ala	Phe 325	Cys	His	Gln	Thr	Glu 330	Thr	Ser	Pro	Lys	Leu 335	Met
Glu	Ile	Gly	Lys 340	Glu	Ile	Val	Lys	Lys 345	Cys	Gly	Gly	Val	Pro 350	Leu	Ala
Ala	Lys	Thr 355	Leu	Gly	Gly	Leu	Leu 360	Arg	Phe	Lys	Arg	Glu 365	Glu	Ser	Glu
Trp	Glu 370	His	Val	Arg	Asp	Ser 375	Glu	Ile	Trp	Asn	Leu 380	Pro	Gln	Asp	Glu
Asn 385	Ser	Val	Leu	Pro	Ala 390	Leu	Arg	Leu	Ser	Tyr 395	His	His	Leu	Pro	Leu 400
Asp	Leu	Arg	Gln	Cys 405	Phe	Ala	Tyr	Суѕ	Ala 410	Val	Phe	Pro	Lys	Asp 415	Thr
Lys	Ile	Glu	Lys 420	Glu	Tyr	Leu	Ile	Ala 425	Leu	Trp	Met	Ala	His 430	Ser	Phe
Leu	Leu	Ser 435	Lys	Gly	Asn	Met	Glu 440	Leu	Glu	Asp	Val	Gly 445	Asn	Glu	Val
Trp	Asn 450	Glu	Leu	Tyr	Leu	Arg 455	Ser	Phe	Phe	Gln	Glu 460	Ile	Glu	Val	Lys
Ser 465	Gly	Lys	Thr	Tyr	Phe 470	Lys	Met	His	Asp	Leu 475	Ile	His	Asp	Leu	Ala 480
Thr	Ser	Met		Ser 485	Ala	Ser	Ala	Ser	Ser 490	Arg	Ser	Ile	Arg	Gln 495	Ile
Asn	Val	Lys	Asp 500	Asp	Glu	Asp	Met	Met 505	Phe	Ile	Val	Thr	Asn 510	Tyr	Lys
Asp	Met	Met 515	Ser	Ile	Gly	Phe	Ser 520	Glu	Val	Val	Ser	Ser 525	Туг	Ser	Pro
Ser	Leu 530	Phe	Lys	Arg	Phe	Val 535	Ser	Leu	Arg	Val	Leu 540	Asn	Leu	Ser	Asn
Ser 545	Glu	Phe	Glu	Gln	Leu 550	Pro	Ser	Ser	Val	Gly 555	Asp	Leu	Val	His	Leu 560

Arg	Tyr	Leu	Asp	Leu	Ser	Gly	Asn	Lys				'25.t Leu		Lys	Arg
	-			565		-		-	570	-				575	,
Leu	Cys	Lys	Leu 580	Gln	Asn	Leu	Gln	Thr 585	Leu	Asp	Leu	Tyr	Asn 590	Cys	Gln
Ser	Leu	Ser 595	Cys	Leu	Pro	Lys	Gln 600	Thr	Ser	Lys	Leu	Cys 605	Ser	Leu	Arg
Asn	Leu 610	Val	Leu	Asp	His	Cys 615	Pro	Leu	Thr	Ser	Met 620	Pro	Pro	Arg	Ile
Gly 625	Leu	Leu	Thr	Cys	Leu 630	Lys	Thr	Leu	Gly	Tyr 635	Phe	Val	Val	Gly	Glu 640
Arg	Lys	Gly	Tyr	Gln 645	Leu	Gly	Glu	Leu	Arg 650	Asn	Leu	Asn	Leu	Arg 655	Gly
Ala	Ile	Ser	Ile 660	Thr	His	Leu	Glu	Arg 665	Val	Lys	Asn	Asp	Met 670	Glu	Ala
Lys	Glu	Ala 675	Asn	Leu	Ser	Ala	Lys 680	Ala	Asn	Leu	His	Ser 685	Leu	Ser	Met
Ser	Trp 690	Asp	Arg	Pro	Asn	Arg 695	Tyr	Glu	Ser	Glu	Glu 700	Val	Lys	Val	Leu
Glu 705	Ala	Leu	Lys	Pro	His 710	Pro	Asn	Leu	Lys	Tyr 715	Leu	Glu	Ile	Ile	Asp 720
Phe	Cys	Gly	Phe	Cys 725	Leu	Pro	Asp	Trp	Met 730	Asn	His	Ser	Val	Leu 735	Lys
Asn	Val	Val	Ser 740	Ile	Leu	Ile	Ser	Gly 745	Суѕ	Glu	Asn	Суѕ	Ser 750	Cys	Leu
Pro	Pro	Phe 755	Gly	Glu	Leu	Pro	Cys 760	Leu	Glu	Ser	Leu	Glu 765	Leu	Gln	Asp
Gly	Ser 770	Val	Glu	Val	Glu	Tyr 775	Val	Glu	Asp	Ser	Gly 780	Phe	Leu	Thr	Arg
Arg 785	Arg	Phe	Pro	Ser	Leu 790	Arg	Lys	Leu	His	Ile 795	Gly	Gly	Phe	Cys	Asn 800
Leu	Lys	Gly	Leu	Gln 805	Arg	Met	Lys	Gly	Ala 810	Glu	Gln	Phe	Pro	Val 815	Leu
Glu	Glu	Met	Lys 820	Ile	Ser	Asp	Cys	Pro 825	Met	Phe	Val	Phe	Pro 830	Thr	Leu
Ser	Ser	Val 835	Lys	Lys	Leu	Glu	Ile 840	Trp	Gly		Ala	845	Ala	Gly	Gly

Leu Ser Ser 850	Ile Ser	Asn Leu 855		Leu Th	hr Ser 860	Leu Lys	Ile	Phe
Ser Asn His 865	Thr Val	Thr Ser 870	Leu Leu		lu Met 75	Phe Lys	Asn	Leu 880
Glu Asn Leu	Ile Tyr 885	Leu Ser	Val Ser	Phe Le 890	eu Glu	Asn Leu	Lys 895	Glu
Leu Pro Thr	Ser Leu 900	Ala Ser	Leu Asr 905		eu Lys	Cys Leu 910	-	Ile
Arg Tyr Cys 915	Tyr Ala	Leu Glu	Ser Leu 920	Pro Gl	lu Glu	Gly Leu 925	Glu	Gly
Leu Ser Ser 930	Leu Thr	Glu Leu 935		Glu H	is Cys 940	Asn Met	Leu	Lys
Cys Leu Pro 945	Glu Gly	Leu Gln 950	His Leu		hr Leu 55	Thr Ser	Leu	Lys 960
Ile Arg Gly	Cys Pro 965	Gln Leu	Ile Lys	Arg Cy 970	ys Glu	Lys Gly	Ile 975	Gly
Glu Asp Trp	His Lys 980	Ile Ser	His Ile 985		sn Val	Asn Ile 990	_	Ile
<210> 11 <211> 2967 <212> DNA <213> Sola:	num bulb	ocastanu	m					
<220> <221> CDS <222> (1).	. (2967)							
<400> 11 atg gcg gaa Met Ala Glu 1								
atc gga gat Ile Gly Asp								
aag ctg tcg Lys Leu Ser 35				-			_	-
cag gag aag Gln Glu Lys 50	_			_	-		_	
ctc aat tct Leu Asn Ser 65					le Leu			
aat gag gca								cca 288

Page 41

Asn	Glu	Ala	Ile	Arg 85	Phe	Glu	Gln	Ser			Gly			His 95	Pro	
					cgt Arg											336
					gca Ala											384
ctt Leu	gaa Glu 130	aaa Lys	att Ile	aca Thr	gag Glu	aga Arg 135	caa Gln	gct Ala	gcc Ala	gct Ala	gct Ala 140	acg Thr	cgt Arg	gaa Glu	aca Thr	432
					gaa Glu 150											480
-					att Ile	_				-		_	_	_	_	528
					ata Ile											576
					ttc Phe											624
					tgt Cys											672
att Ile 225	aag Lys	aca Thr	att Ile	ata Ile	gga Gly 230	aat Asn	att Ile	gaa Glu	aga Arg	agt Ser 235	tct Ser	cct Pro	cat His	gtt Val	gag Glu 240	720
gac Asp	ttg Leu	gct Ala	tca Ser	ttt Phe 245	cag Gln	aag Lys	aag Lys	ctc Leu	cag Gln 250	gag Glu	tta Leu	ttg Leu	aat Asn	gga Gly 255	aaa Lys	768
					tta Leu											816
tgg Trp	gct Ala	aag Lys 275	tta Leu	aga Arg	gca Ala	Val	tta Leu 280	act Thr	gtt Val	gga Gly	gca Ala	aga Arg 285	ggt Gly	gct Ala	tct Ser	864
					cgt Arg											912
_					ttg Leu 310			_				_	_			960
					gca Ala											1008
					aag Lys											1056
					ctt Leu											1104

	gaa Glu 370															1	152
	gaa Glu															1	200
cca Pro	ctt Leu	gat Asp	ttg Leu	aga Arg 405	caa Gln	tgc Cys	ttt Phe	gcg Ala	tat Tyr 410	tgt Cys	gca Ala	gta Val	ttc Phe	cca Pro 415	aag Lys	. 1	248
	acc Thr															1	296
	ttt Phe															1	344
	gta Val 450															1	392
	aaa Lys															1	440
	gct Ala															1	488
	ata Ile															1	536
-	gtg Val							_		_		_		_	_	1	584
	agg Arg 530															1	632
	att Ile															1	680
	ttc Phe															1	728
	ctt Leu															1	776
	agt Ser															1	824
	act Thr 610															1	872
	ggt Gly															1	920
ctg	aaa	aac	cta	aat	ctc	tgc	ggc	tca	att	tca	atc	aca	cac	ctt	gag	1	968

Leu	Lys	Asn	Leu		Leu	Cys	Gly	Ser	Ile	5.ST Ile				Glu	
-				645					650				655		
					acg Thr										2016
					agc Ser										2064
					gtt Val										2112
					gag Glu 710										2160
					tca Ser										2208
					tgc Cys										2256
					gag Glu										2304
					gtc Val										2352
					ctt Leu 790										2400
					gga Gly										2448
					cct Pro										2496
					cac His										2544
					act Thr										2592
_					cca Pro 870	-		-		_					2640
_		-	_		ttt Phe	-		_			-	_			2688
					aat Asn										2736
-				_	ttc Phe		-			-					2784

Page 44

			aag atg cta Lys Met Leu 940		
		Thr Ala Leu	aca aat tta Thr Asn Leu 955		
			aag gaa ata Lys Glu Ile 970		Trp
	-	_	gat att cat Asp Ile His	-	2967
<210> 12 <211> 988 <212> PRT <213> Solar	num bulbocas	tanum			
<400> 12					
Met Ala Glu 1	Ala Phe Leu 5	Gln Val Leu	Leu Glu Asn 10	Leu Thr Ser 15	? Phe
Ile Gly Asp	Lys Leu Val 20	Leu Ile Phe 25	Gly Phe Glu	Lys Glu Cys	s Glu
Lys Leu Ser 35	Ser Val Phe	Ser Thr Ile	Gln Ala Val	Leu Gln Asp 45	o Ala
Gln Glu Lys 50	Gln Leu Lys	Asp Lys Ala 55	Ile Glu Asn 60	Trp Leu Gli	n Lys
Leu Asn Ser 65	Ala Ala Tyr 70	Glu Val Asp	Asp Ile Leu 75	Gly Glu Cys	s Lys 80
Asn Glu Ala	Ile Arg Phe 85	Glu Gln Ser	Arg Leu Gly 90	Phe Tyr His	s Pro
Gly Ile Ile	Asn Phe Arg	His Lys Ile 105	Gly Arg Arg	Met Lys Glu	ı Ile
Met Glu Lys 115	Leu Asp Ala	Ile Ser Glu 120	Glu Arg Arg	Lys Phe His	s Phe
Leu Glu Lys 130	Ile Thr Glu	Arg Gln Ala 135	Ala Ala Ala 140	Thr Arg Glu	1 Thr
Gly Phe Val 145	Leu Thr Glu 150	Pro Lys Val	Tyr Gly Arg 155	Asp Lys Glu	1 Glu 160
Asp Glu Ile	Val Lys Ile 165	Leu Ile Asn	Asn Val Asn 170	Val Ala Glu 17	

Page 45

Leu Pro Val Phe Pro Ile Ile Gly Met Gly Gly Leu Gly Lys Thr Thr

Leu	Ala	Gln 195	Met	Ile	Phe	Asn	Asp 200	Glu	Arg	Val	Thr	Lys 205	His	Phe	Asn
Pro	Lys 210	Ile	Trp	Val	Cys	Val 215	Ser	Asp	Asp	Phe	Asp 220	Glu	Lys	Arg	Leu
Ile 225	Lys	Thr	Ile	Ile	Gly 230	Asn	Ile	Glu	Arg	Ser 235	Ser	Pro	His	Val	Glu 240
Asp	Leu	Ala	Ser	Phe 245	Gln	Lys	Lys	Leu	Gln 250	Glu	Leu	Leu	Asn	Gly 255	Lys
Arg	Tyr	Leu	Leu 260	Val	Leu	Asp	Asp	Val 265	Trp	Asn	Asp	Asp	Leu 270	Glu	Lys
Trp	Ala	Lys 275	Leu	Arg	Ala	Val	Leu 280	Thr	Val	Gly	Ala	Arg 285	Gly	Ala	Ser
Ile	Leu 290	Ala	Thr	Thr	Arg	Leu 295	Glu	Lys	Val	Gly	Ser 300	Ile	Met	Gly	Thr
Leu 305	Gln	Pro	Tyr	His	Leu 310	Ser	Asn	Leu	Ser	Pro 315	His	Asp	Ser	Leu	Leu 320
Leu	Phe	Met	Gln	Arg 325	Ala	Phe	Gly	Gln	Gln 330	Lys	Glu	Ala	Asn	Pro 335	Asn
Leu	Val	Ala	Ile 340	Gly	Lys	Glu	Ile	Val 345	Lys	Lys	Cys	Gly	Gly 350	Val	Pro
Leu	Ala	Ala 355	Lys	Thr	Leu	Gly	Gly 360	Leu	Leu	Arg	Phe	Lys 365	Arg	Glu	Glu
Ser	Glu 370	Trp	Glu	His	Val	Arg 375	Asp	Asn	Glu	Ile	Trp 380	Ser	Leu	Pro	Gln
Asp 385	Glu	Ser	Ser	Ile	Leu 390	Pro	Ala	Leu	Arg	Leu 395	Ser	Tyr	His	His	Leu 400
Pro	Leu	Asp	Leu	Arg 405	Gln	Cys	Phe	Ala	Tyr 410	Cys	Ala	Val	Phe	Pro 415	Lys
Asp	Thr	Lys	Met 420	Ile	Lys	Glu	Asn	Leu 425	Ile	Thr	Leu	Trp	Met 430	Ala	His
Gly	Phe	Leu 435	Leu	Ser	Lys	Gly	Asn 440	Leu	Glu	Leu	Glu	Asp 445	Val	Gly	Asn
Glu	Val 450	Trp	Asn	Glu	Leu	Tyr 455	Leu	Arg	Ser	Phe	Phe 460	Gln	Glu	Ile	Glu

Ala 465	Lys	Ser	Gly	Asn	Thr 470	Tyr	Phe	Lys	Ile	His 475	Asp	Leu	Ile	His	Asp 480
Leu	Ala	Thr	Ser	Leu 485	Phe	Ser	Ala	Ser	Ala 490	Ser	Суѕ	Gly	Asn	Ile 495	Arg
Glu	Ile	Asn	Val 500	Lys	Asp	Tyr	Lys	His 505	Thr	Val	Ser	Ile	Gly 510	Phe	Ala
Ala	Val	Val 515	Ser	Ser	Tyr	Ser	Pro 520	Ser	Leu	Leu	Lys	Lys 525	Phe	Val	Ser
Leu	Arg 530	Val	Leu	Asn	Leu	Ser 535	Tyr	Ser	Lys	Leu	Glu 540	Gln	Leu	Pro	Ser
Ser 545	Ile	Gly	Asp	Leu	Leu 550	His	Leu	Arg	Tyr	Leu 555	Asp	Leu	Ser	Cys	Asn 560
Asn	Phe	Arg	Ser	Leu 565	Pro	Glu	Arg	Leu	Cys 570	Lys	Leu	Gln	Asn	Leu 575	Gln
Thr	Leu	Asp	Val 580	His	Asn	Cys	Tyr	Ser 585	Leu	Asn	Cys	Leu	Pro 590	Lys	Gln
Thr	Ser	Lys 595	Leu	Ser	Ser	Leu	Arg 600	His	Leu	Val	Val	Asp 605	Gly	Cys	Pro
Leu	Thr 610	Ser	Thr	Pro	Pro	Arg 615	Ile	Gly	Leu	Leu	Thr 620	Cys	Leu	Lys	Thr
Leu 625	Gly	Phe	Phe	Ile	Val 630	Gly	Ser	Lys	Lys	Gly 635	Tyr	Gln	Leu	Gly	Glu 640
Leu	Lys	Asn	Leu	Asn 645	Leu	Cys	Gly	Ser	Ile 650	Ser	Ile	Thr	His	Leu 655	Glu
Arg	Val	Lys	Asn 660	Asp	Thr	Asp	Ala	Glu 665	Ala	Asn	Leu	Ser	Ala 670	Lys	Ala
Asn	Leu	Gln 675	Ser	Leu	Ser	Met	Ser 680	Trp	Asp	Asn	Asp	Gly 685	Pro	Asn	Arg
Tyr	Glu 690	Ser	Lys	Glu	Val	Lys 695	Val	Leu	Glu	Ala	Leu 700	Lys	Pro	His	Pro
Asn 705	Leu	Lys	Tyr	Leu	Glu 710	Ile	Ile	Ala	Phe	Gly 715	Gly	Phe	Arg	Phe	Pro 720
Ser	Trp	Ile	Asn	His 725	Ser	Val	Leu	Glu	Lys 730	Val	Ile	Ser	Val	Arg 735	Ile
Lys	Ser	Cys	Lys	Asn	Cys	Leu	Cys	Leu	Pro		Phe ge 4	_	Glu	Leu	Pro

Cys Leu Glu Asn Leu Glu Leu Gln Asn Gly Ser Ala Glu Val Glu Tyr 755 760 765

Val Glu Glu Asp Asp Val His Ser Arg Phe Ser Thr Arg Arg Ser Phe 770 780

Pro Ser Leu Lys Lys Leu Arg Ile Trp Phe Phe Arg Ser Leu Lys Gly 785 790 795

Leu Met Lys Glu Glu Glu Glu Lys Phe Pro Met Leu Glu Glu Met 805 810 815

Ala Ile Leu Tyr Cys Pro Leu Phe Val Phe Pro Thr Leu Ser Ser Val 820 825 830

Lys Lys Leu Glu Val His Gly Asn Thr Asn Thr Arg Gly Leu Ser Ser 835 840 845

Ile Ser Asn Leu Ser Thr Leu Thr Ser Leu Arg Ile Gly Ala Asn Tyr 850 855 860

Arg Ala Thr Ser Leu Pro Glu Glu Met Phe Thr Ser Leu Thr Asn Leu 865 870 875 880

Glu Phe Leu Ser Phe Phe Asp Phe Lys Asn Leu Lys Asp Leu Pro Thr 885 890 895

Ser Leu Thr Ser Leu Asn Ala Leu Lys Arg Leu Gln Ile Glu Ser Cys 900 905 910

Asp Ser Leu Glu Ser Phe Pro Glu Gln Gly Leu Glu Gly Leu Thr Ser 915 920 925

Leu Thr Gln Leu Phe Val Lys Tyr Cys Lys Met Leu Lys Cys Leu Pro 930 935 940

Glu Gly Leu Gln His Leu Thr Ala Leu Thr Asn Leu Gly Val Ser Gly 945 955 960

Cys Pro Glu Val Glu Lys Arg Cys Asp Lys Glu Ile Gly Glu Asp Trp 965 970 975

His Lys Ile Ala His Ile Pro Asn Leu Asp Ile His

<210> 13

<211> 1920

<212> DNA

<213> Solanum bulbocastanum

<220>

<221>	CDS
<222>	(1)(1920)

atg gct gaa gct ttc att caa gtt gtg cta gac aat ctc act tct ttc 48 Met Ala Glu Ala Phe Ile Gln Val Val Leu Asp Asn Leu Thr Ser Phe ctc aaa ggg gaa ctt gta ttg ctt ttc ggt ttt caa gat gag ttc caa 96 Leu Lys Gly Glu Leu Val Leu Phe Gly Phe Gln Asp Glu Phe Gln agg ctt tca agc atg ttt tct aca atc caa gcc gtc ctt gaa gat qct 144 Arg Leu Ser Ser Met Phe Ser Thr Ile Gln Ala Val Leu Glu Asp Ala 40 caa gag aag caa ctc aac gac aag cct cta gaa aat tgg ttg caa aaa 192 Gln Glu Lys Gln Leu Asn Asp Lys Pro Leu Glu Asn Trp Leu Gln Lys ctc aat gct gct aca tat gaa gtc gat gac atc ttg gat gaa tat aaa 240 Leu Asn Ala Ala Thr Tyr Glu Val Asp Asp Ile Leu Asp Glu Tyr Lys act aag gcc aca aga ttc ttg ctg tct gaa tat ggc cgt tat cat cca 288 Thr Lys Ala Thr Arg Phe Leu Leu Ser Glu Tyr Gly Arg Tyr His Pro aag gtt atc cct ttc cgt cac aag gtt ggg aaa agg atg gac caa gtg Lys Val Ile Pro Phe Arg His Lys Val Gly Lys Arg Met Asp Gln Val 336 atg aaa aaa ctg aat gca att gct gag gaa cga aag aat ttt cat ttg 384 Met Lys Lys Leu Asn Ala Ile Ala Glu Glu Arg Lys Asn Phe His Leu caa gaa aag att ata gag aga caa gct gct aca cgg gaa aca ggt tct 432 Gln Glu Lys Ile Ile Glu Arg Gln Ala Ala Thr Arg Glu Thr Gly Ser 135 140 gtg tta act gaa tca caa gtt tat gga agg gac aaa gaa aaa gat gag 480 Val Leu Thr Glu Ser Gln Val Tyr Gly Arg Asp Lys Glu Lys Asp Glu 145 150 155 ata gtg aaa atc cta aca aac act gct agt gat gcc caa aaa ctc tca Ile Val Lys Ile Leu Thr Asn Thr Ala Ser Asp Ala Gln Lys Leu Ser 165 170 gtc ctc cca ata ctt ggt atg ggg gga cta gga aag acg act ctt tcc 576 Val Leu Pro Ile Leu Gly Met Gly Gly Leu Gly Lys Thr Thr Leu Ser caa atg gtc ttc aat gat cag aga gta act gag cgt ttc tat ccc aaa 624 Gln Met Val Phe Asn Asp Gln Arg Val Thr Glu Arg Phe Tyr Pro Lys 200 ata tgg att tgc gtc tcg gat gat ttt aat gag aag agg ttg ata aag 672 Ile Trp Ile Cys Val Ser Asp Asp Phe Asn Glu Lys Arg Leu Ile Lys gca ata gta gaa tct att gaa ggg aag tcc ctc agt gac atg gac ttg 720 Ala Ile Val Glu Ser Ile Glu Gly Lys Ser Leu Ser Asp Met Asp Leu 230 235 gct cca ctt caa aag aag ctt caa gag ttg ctg aat gga aaa aga tac 768 Ala Pro Leu Gln Lys Lys Leu Gln Glu Leu Leu Asn Gly Lys Arg Tyr 250 ttc ctt gtc tta gat gat gtt tgg aat gaa gat caa cat aag tgg gct 816 Phe Leu Val Leu Asp Asp Val Trp Asn Glu Asp Gln His Lys Trp Ala

					ttg Leu											864
					gaa Glu											912
					aat Asn 310											960
					gga Gly											1008
					att Ile											1056
gcc Ala	aag Lys	act Thr 355	ctt Leu	gga Gly	ggt Gly	att Ile	ttg Leu 360	cgc Arg	ttc Phe	aag Lys	aga Arg	gaa Glu 365	gaa Glu	aga Arg	gaa Glu	1104
					gac Asp											1152
					gcc Ala 390											1200
					ttt Phe											1248
					aat Asn											1296
		_			aat Asn	_				_	_			_	_	1344
					ttg Leu							Ile				1392
					ttc Phe 470											1440
					gca Ala											1488
					gga Gly											1536
					cct Pro											1584
				-	aac Asn	_										1632

													25. t					
(								tac Tyr									1680	
		-	_				-	tta Leu	_	_				_	_		1728	
								tct Ser									1776	
								aat Asn 600									1824	
								gga Gly									1872	j
S								aga Arg								taa	1920	
<	(210) (211) (212) (213) (400)	> 6 2> E 3> S	l4 539 PRT Solar	num l	oulbo	ocast	canur	n										
	1et			Ala	Phe 5	Ile	Gln	Val	Val	Leu 10	Asp	Asn	Leu	Thr	Ser 15	Phe		
Ι	₋eu	Lys	Gly	Glu 20	Leu	Val	Leu	Leu	Phe 25	Gly	Phe	Gln	Asp	Glu 30	Phe	Gln		
Į	Arg	Leu	Ser 35	Ser	Met	Phe	Ser	Thr 40	Ile	Gln	Ala	Val	Leu 45	Glu	Asp	Ala		
C	Sln	Glu 50	Lys	Gln	Leu	Asn	Asp 55	Lys	Pro	Leu	Glu	Asn 60	Trp	Leu	Gln	Lys		
	eu 55	Asn	Ala	Ala	Thr	Tyr 70	Glu	Val	Asp	Asp	Ile 75	Leu	Asp	Glu	Tyr	Lys 80		
1	hr:	Lys	Ala	Thr	Arg 85	Phe	Leu	Leu	Ser	Glu 90	Tyr	Gly	Arg	Tyr	His 95	Pro		
Ι	ys	Val	Ile	Pro 100	Phe	Arg	His	Lys	Val 105	Gly	Lys	Arg	Met	Asp 110	Gln	Val		
M	let	Lys	Lys 115	Leu	Asn	Ala	Ile	Ala 120		Glu	Arg	Lys	Asn 125	Phe	His	Leu		

Gln Glu Lys Ile Ile Glu Arg Gln Ala Ala Thr Arg Glu Thr Gly Ser  $130\,$ 

Val Leu Thr Glu Ser Gln Val Tyr Gly Arg Asp Lys Glu Lys Asp Glu 145 150 155 160

Page 51

Ile	Val	Lys	Ile	Leu 165	Thr	Asn	Thr	Ala	Ser 170	Asp	Ala	Gln	Lys	Leu 175	Ser
Val	Leu	Pro	Ile 180	Leu	Gly	Met	Gly	Gly 185	Leu	Gly	Lys	Thr	Thr 190	Leu	Ser
Gln	Met	Val 195	Phe	Asn	Asp	Gln	Arg 200	Val	Thr	Glu	Arg	Phe 205	Tyr	Pro	Lys
Ile	Trp 210	Ile	Cys	Val	Ser	Asp 215	Asp	Phe	Asn	Glu	Lys 220	Arg	Leu	Ile	Lys
Ala 225	Ile	Val	Glu	Ser	Ile 230	Glu	Gly	Lys	Ser	Leu 235	Ser	Asp	Met	Asp	Leu 240
Ala	Pro	Leu	Gln	Lys 245	Lys	Leu	Gln	Glu	Leu 250	Leu	Asn	Gly	Lys	Arg 255	Tyr
Phe	Leu	Val	Leu 260	Asp	Asp	Val	Trp	Asn 265	Glu	Asp	Gln	His	Lys 270	Trp	Ala
Asn	Leu	Arg 275	Ala	Val	Leu	Lys	Val 280	Gly	Ala	Ser	Gly	Ala 285	Phe	Val	Leu
Thr	Thr 290	Thr	Arg	Leu	Glu	Lys 295	Val	Gly	Ser	Ile	Met 300	Gly	Thr	Leu	Gln
Pro 305	Tyr	Glu	Leu	Ser	Asn 310	Leu	Ser	Pro	Glu	Asp 315	Cys	Trp	Phe	Leu	Phe 320
Met	Gln	Arg	Ala	Phe 325	Gly	His	Gln	Glu	Glu 330	Ile	Asn	Pro	Asn	Leu 335	Val
Ala	Ile	Gly	Lys 340	Glu	Ile	Val	Lys	Lys 345	Cys	Gly	Gly	Val	Pro 350	Leu	Ala
Ala	Lys	Thr 355	Leu	Gly	Gly	Ile	Leu 360	Arg	Phe	Lys	Arg	Glu 365	Glu	Arg	Glu
Trp	Glu 370	His	Val	Arg	Asp	Ser 375	Pro	Ile	Trp	Asn	Leu 380	Pro	Gln	Asp	Glu
Ser 385	Ser	Ile	Leu	Pro	Ala 390	Leu	Arg	Leu	Ser	Tyr 395	His	His	Leu	Pro	Leu 400
Asp	Leu	Arg	Gln	Cys 405	Phe	Val	Tyr	Cys	Ala 410	Val	Phe	Pro	Lys	Asp 415	Thr
Lys	Met	Ala	Lys 420	Glu	Asn	Leu	Ile	Ala 425	Phe	Trp	Met	Ala	His 430	Gly	Phe

Leu	Leu	Ser 435	Lys	Gly	Asn	Leu	Glu 440	Leu		F023 Asp				Glu	Val	
Trp	Asn 450	Glu	Leu	Tyr	Leu	Arg 455	Ser	Phe	Phe	Gln	Glu 460	Ile	Glu	Val	Glu	
Ser 465	Gly	Lys	Thr	Tyr	Phe 470	Lys	Met	His	Asp	Leu 475	Ile	His	Asp	Leu	Ala 480	
Thr	Ser	Leu	Phe	Ser 485	Ala	Asn	Thr	Ser	Ser 490	Ser	Asn	Ile	Arg	Glu 495	Ile	
Asn	Ala	Asn	Tyr 500	Asp	Gly	Tyr	Met	Met 505	Ser	Ile	Gly	Phe	Ala 510	Glu	Val	
Val	Ser	Ser 515	Tyr	Ser	Pro	Ser	Leu 520	Leu	Gln	Lys	Phe	Val 525	Ser	Leu	Arg	
Val	Leu 530	Asn	Leu	Arg	Asn	Ser 535	Asn	Leu	Asn	Gln	Leu 540	Pro	Ser	Ser	Ile	
Gly 545	Asp	Leu	Val	His	Leu 550	Arg	Tyr	Leu	Asp	Leu 555	Ser	Gly	Asn	Val	Arg 560	
Ile	Arg	Ser	Leu	Pro 565	Arg	Arg	Leu	Cys	Lys 570	Leu	Gln	Asn	Leu	Gln 575	Thr	
Leu	Asp	Leu	His 580	Tyr	Cys	Asp	Ser	Leu 585	Ser	Суѕ	Leu	Pro	Lys 590	Gln	Thr	
Ser	Lys	Leu 595	Gly	Ser	Leu	Arg	Asn 600	Leu	Leu	Leu	Asp	Gly 605	Cys	Ser	Leu	
Thr	Ser 610	Thr	Pro	Pro	Arg	Ile 615	Gly	Leu	Leu	Thr	Cys 620	Leu	Lys	Ser	Leu	
Ser 625	Cys	Phe	Val	Ile	Gly 630	Lys	Arg	Lys	Val	Ile 635	Asn	Leu	Val	Asn		
<210 <211 <211 <211	1> 8 2> [	L5 879 DNA Solar	num k	oulbo	ocast	canur	n	-								
<220 <221 <221	1> (	CDS (1)	. (879	9)												
	0> 1 tgg Trp															48
	gga Gly															96
										_	-	_				

Page 53

					caa Gln											144
					acg Thr											192
aca Thr 65	gta Val	act Thr	gag Glu	cat His	ttg Leu 70	tat Tyr	ccg Pro	aaa Lys	ata Ile	tgg Trp 75	att Ile	tgt Cys	gtc Val	tcc Ser	aat Asn 80	240
					agg Arg											288
					ggt Gly											336
					atg Met											384
					ccc Pro											432
					ttt Phe 150											480
gga Gly	gaa Glu	gag Glu	caa Gln	ttc Phe 165	cct Pro	gtg Val	ctt Leu	gaa Glu	gag Glu 170	atg Met	gag Glu	att Ile	aaa Lys	tgg Trp 175	tgc Cys	528
					ccg Pro											576
					gat Asp											624
agg Arg	gct Ala 210	ctt Leu	act Thr	tcc Ser	ctc Leu	aat Asn 215	att Ile	aac Asn	ttt Phe	aac Asn	aaa Lys 220	gaa Glu	gct Ala	act Thr	tca Ser	672
-	-	~ 1	~ 1		ttc Phe 230	-	~	-	- 1	_	-	_		_	_	720
					aat Asn											768
					agt Ser											816
-			_	_	ggg Gly										-	864
	gtc Val 290		gac Asp	tga												879

- <210> 16 <211> 292
- <212> PRT
- <213> Solanum bulbocastanum

<400> 16

Met Trp Thr Leu Leu Gly Pro Gly Ser Val Leu Thr Glu Pro Gln Val

Tyr Gly Arg Asp Lys Glu Lys Asp Glu Ile Val Lys Ile Leu Ile Asn

Asn Val Ser Asp Ala Gln Glu Val Ser Val Leu Pro Ile Val Gly Met

Gly Gly Leu Gly Lys Thr Thr Leu Ala Gln Met Val Phe Asn Asp Gln

Thr Val Thr Glu His Leu Tyr Pro Lys Ile Trp Ile Cys Val Ser Asn

Asp Phe Asp Glu Lys Arg Leu Ile Lys Ala Ile Val Glu Ser Ile Glu

Gly Arg Pro Leu Gly Glu Met Asp Leu Ala Pro Leu Gln Lys Lys

Leu Gln Glu Leu Arg Met Trp Ser Met Leu Lys Lys Trp Ile Leu Met

Phe Ile Leu Asp Phe Pro Thr Arg Ile Arg Phe Pro Ser Leu Arg Lys

Leu Asp Ile Trp Asp Phe Gly Ser Leu Lys Gly Leu Leu Lys Lys Glu

Gly Glu Glu Gln Phe Pro Val Leu Glu Glu Met Glu Ile Lys Trp Cys

Pro Met Phe Val Ile Pro Thr Leu Ser Ser Val Lys Lys Leu Val Val 185

Arg Gly Asp Lys Ser Asp Ala Ile Gly Phe Ser Ser Ile Ser Asn Leu

Arg Ala Leu Thr Ser Leu Asn Ile Asn Phe Asn Lys Glu Ala Thr Ser

Leu Pro Glu Glu Met Phe Lys Ser Leu Ala Asn Leu Lys Tyr Leu Lys

Ile Ser Ser Phe Arg Asn Leu Lys Glu Leu Pro Thr Ser Leu Ala Ser 245 250

									WAL	CF UZ S	55.51	25.0	.xc			
Leu	Asn	Ala	Leu 260	Gln	Ser	Leu	Thr	Ile 265	Glu	His	Cys	Asp	Ala 270	Leu	Glu	
Ser	Leu	Pro 275	Glu	Glu	Gly	Val	Lys 280	Gly	Leu	Thr	Ser	Leu 285	Thr	Glu	Leu	
Ser	Val 290	Gln	Asp													
<210 <213 <213 <213	l> : 2> i	17 1362 DNA Solar	num l	oulbo	ocast	canur	n									
<220 <221 <222	L> (	CDS (1).	. (136	52)												
	gct			ttc Phe 5												48
				ctt Leu												96
				atg Met												144
cag Gln	gag Glu 50	aag Lys	caa Gln	ctc Leu	aac Asn	aac Asn 55	aag Lys	cct Pro	cta Leu	gaa Glu	aat Asn 60	tgg Trp	ttg Leu	caa Gln	aaa Lys	192
				aca Thr												240
				aga Arg 85												288
aag Lys	gtt Val	atc Ile	cct Pro 100	ttc Phe	cgt Arg	cac His	aag Lys	gtc Val 105	Gly	aaa Lys	agg Arg	atg Met	gac Asp 110	caa Gln	gtg Val	336
Met	Lys	Lys	Leu	aag Lys	Ala	Ile	Ala	Glu	gaa Glu	Arg	Lys	Asn	Phe	cat His	ttg Leu	384

 $1\bar{1}5$ 

120

150

165

cac gaa aaa att gta gag aga caa gct gtt aga cgg gaa aca ggt tct

His Glu Lys Ile Val Glu Arg Gln Ala Val Arg Arg Glu Thr Gly Ser

gta tta acc gaa ccg cag gtt tat gga aga gac aaa gag aaa gat gag Val Leu Thr Glu Pro Gln Val Tyr Gly Arg Asp Lys Glu Lys Asp Glu

ata gtg aaa atc cta ata aac aat gtt agt gat gcc caa cac ctt tca

Ile Val Lys Ile Leu Ile Asn Asn Val Ser Asp Ala Gln His Leu Ser

gtc ctc cca ata ctt ggt atg ggg gga tta gga aaa acg act ctt gcc Val Leu Pro Ile Leu Gly Met Gly Gly Leu Gly Lys Thr Thr Leu Ala

Page 56

155

170

432

480

528

576

	-	-			-	cag Gln	_	_							624
						gaa Glu 215									672
						gaa Glu									720
-	-				_	aag Lys		-	_	_	_			_	768
						gat Asp									816
						ttg Leu									864
						gaa Glu 295									912
						aat Asn									960
						gga Gly									1008
	-			_		att Ile				_					1056
						ggt Gly									1104
-		-			_	gac Asp 375	-	-				_		_	1152
	_			-		gcc Ala	_			_					1200
	_	-			_	ttt Phe			_				_	_	1248
-		_	-		-	aag Lys						_			1296
						aac Asn	_				-		 _		1344
		aaa Lys	-		tag										1362

									WAR	RF023	85.ST	25.t	xt		
<210 <210 <210 <210	l> 2>	18 453 PRT Sola:	num 1	oulbo	ocast	tanur	n								
<400	>	18													
Met 1	Ala	Glu	Ala	Phe 5	Ile	Gln	Val	Leu	Leu 10	Asp	Asn	Leu	Thr	Ser 15	Phe
Leu	Lys	Gly	Glu 20	Leu	Ala	Leu	Leu	Phe 25	Gly	Phe	Gln	Asp	Glu 30	Phe	Gln
Arg	Leu	Ser 35	Ser	Met	Phe	Ser	Thr 40	Ile	Gln	Ala	Val	Leu 45	Glu	Asp	Ala
Gln	Glu 50	Lys	Gln	Leu	Asn	Asn 55	Lys	Pro	Leu	Glu	Asn 60	Trp	Leu	Gln	Lys
Leu 65	Asn	Ala	Ala	Thr	Tyr 70	Glu	Val	Asp	Asp	Ile 75	Leu	Asp	Glu	Tyr	Lys 80
Thr	Lys	Ala	Thr	Arg 85	Phe	Ser	Gln	Ser	Glu 90	Tyr	Gly	Arg	Tyr	His 95	Pro
Lys	Val	Ile	Pro 100	Phe	Arg	His	Lys	Val 105	Gly	Lys	Arg	Met	Asp 110	Gln	Val
Met	Lys	Lys 115	Leu	Lys	Ala	Ile	Ala 120	Glu	Glu	Arg	Lys	Asn 125	Phe	His	Leu
His	Glu 130	Lys	Ile	Val	Glu	Arg 135	Gln	Ala	Val	Arg	Arg 140	Glu	Thr	Gly	Ser
Val 145	Leu	Thr	Glu	Pro	Gln 150	Val	Tyr	Gly	Arg	Asp 155	Lys	Glu	Lys	Asp	Glu 160
Ile	Val	Lys	Ile	Leu 165	Ile	Asn	Asn	Val	Ser 170	Asp	Ala	Gln	His	Leu 175	Ser
Val	Leu	Pro	Ile 180	Leu	Gly	Met	Gly	Gly 185	Leu	Gly	Lys	Thr	Thr 190	Leu	Ala
Gln	Met	Val 195	Phe	Asn	Asp	Gln	Arg 200	Val	Thr	Glu	His	Phe 205	His	Ser	Lys
Ile	Trp 210	Ile	Cys	Val	Ser	Glu 215	Asp	Phe	Asp	Glu	Lys 220	Arg	Leu	Ile	Lys
Ala 225	Ile	Val	Glu	Ser	Ile 230	Glu	Gly	Arg	Pro	Leu 235	Leu	Gly	Glu	Met	Asp 240

Leu Ala Pro Leu Gl<br/>n Lys Lys Leu Gl<br/>n Glu Leu Leu As<br/>n Gly Lys Arg245  $\phantom{0}$  250  $\phantom{0}$  255

Page 58

Tyr	Leu	Leu	Val 260	Leu	Asp	Asp	Val	Trp 265	Asn	Glu	Asp	Gln	Gln 270	Lys	Trp	
Ala	Asn	Leu 275	Arg	Ala	Val	Leu	Lys 280	Val	Gly	Ala	Ser	Gly 285	Ala	Ser	Val	
Leu	Thr 290	Thr	Thr	Arg	Leu	Glu 295	Lys	Val	Gly	Ser	Ile 300	Met	Gly	Thr	Leu	
Gln 305	Pro	Tyr	Glu	Leu	Ser 310	Asn	Leu	Ser	Gln	Glu 315	Asp	Cys	Trp	Leu	Leu 320	
Phe	Met	Gln	Arg	Ala 325	Phe	Gly	His	Gln	Glu 330	Glu	Ile	Asn	Pro	Asn 335	Leu	
Val	Ala	Ile	Gly 340	Lys	Glu	Ile	Val	Lys 345	Lys	Ser	Gly	Gly	Val 350	Pro	Leu	
Ala	Ala	Lys 355	Thr	Leu	Gly	Gly	Ile 360	Leu	Cys	Phe	Lys	Arg 365	Glu	Glu	Arg	
Ala	Trp 370	Glu	His	Val	Arg	Asp 375	Ser	Pro	Ile	Trp	Asn 380	Leu	Pro	Gln	Asp	
Glu 385	Ser	Ser	Ile	Leu	Pro 390	Ala	Leu	Arg	Leu	Ser 395	Tyr	His	Gln	Leu	Pro 400	
Leu	Asp	Leu	Lys	Gln 405	Cys	Phe	Ala	Tyr	Cys 410	Ala	Val	Phe	Pro	Lys 415	Asp	
Ala	Lys	Met	Glu 420	Lys	Glu	Lys	Leu	Ile 425	Ser	Leu	Trp	Met	Ala 430	His	Gly	
Phe	Leu	Leu 435	Ser	Lys	Gly	Asn	Met 440	Glu	Leu	Glu	Asp	Val 445	Gly	Asp	Glu	
	Trp 450		Glu	Leu												
<210 <211 <212 <213	L> 3 2> I	19 3096 DNA Solar	num k	oulbo	ocast	anum	n									
<220 <221 <222	L> (	CDS (1)	. (309	96)												
	tgg				ttt Phe											48
cct	tcc	tct	ctt	act	cca	cag	gat	ttc	agc		aga ige 5	-	ttt	aaa	ttg	96

Pro	Ser	Ser	Leu 20	Thr	Pro	Gln	Asp	Phe 25			Arg			Lys	Leu	
aag Lys	aga Arg	aat Asn 35	gtg Val	aag Lys	att Ile	tca Ser	aga Arg 40	aat Asn	ttt Phe	gaa Glu	ttt Phe	gct Ala 45	atc Ile	agt Ser	tgt Cys	144
					gct Ala											192
					ctt Leu 70											240
					ggt Gly											288
					cgt Arg											336
					aga Arg											384
					att Ile											432
					ttt Phe 150											480
					gtc Val											528
					gtc Val											576
					aaa Lys											624
				Ala	caa Gln	Glu	Lys			Lys						672
					ctc Leu 230											720
					act Thr											768
					cgg Arg											816
					atg Met											864
					gat Asp											912

Page 60

cgg Arg 305	caa Gln	aca Thr	ggt Gly	ttt Phe	gtt Val 310	tta Leu	act Thr	gag Glu	cca Pro	aaa Lys 315	gtt Val	tat Tyr	gga Gly	aag Lys	gaa Glu 320	960
		gag Glu														1008
		gaa Glu														1056
aag Lys	acg Thr	act Thr 355	cta Leu	gcc Ala	caa Gln	atg Met	gtc Val 360	ttc Phe	aat Asn	gat Asp	caa Gln	aga Arg 365	att Ile	act Thr	gag Glu	1104
		aat Asn														1152
		ttg Leu														1200
		atg Met														1248
		aaa Lys														1296
		aag Lys 435		-			-	-	_	_	_			_	_	1344
		tca Ser														1392
		act Thr														1440
		ttg Leu														1488
_	_	aaa Lys	-		~ 3	- 1	~ -	_	~ -			_	_	_		1536
ggt Gly	gtg Val	cct Pro 515	cta Leu	gca Ala	gcc Ala	aaa Lys	act Thr 520	ctt Leu	gga Gly	ggc Gly	ctt Leu	tta Leu 525	cgc Arg	ttc Phe	aag Lys	1584
		gaa Glu														1632
		caa Gln														1680
		ctt Leu														1728
ttc	cca	aag	gac	acc	aaa	ata	gaa	aag	gaa		ctc		gct	ctc	tgg	1776

Page 61

Phe	Pro	Lys	Asp 580	Thr	Lys	Ile	Glu	Lys 585			5.ST Leu			Leu	Trp	
					ctt Leu											1824
					tgg Trp											1872
					tct Ser 630											1920
					gat Asp											1968
					att Ile											2016
					ggc Gly											2064
					cgt Arg											2112
					gag Glu 710											2160
		_			agc Ser	_	-		_	_			_		-	2208
					gtg Val											2256
					att Ile											2304
atg Met	aat Asn 770	cac His	tca Ser	gtt Val	ttg Leu	aaa Lys 775	aat Asn	gtt Val	gtc Val	tct Ser	att Ile 780	cta Leu	att Ile	agc Ser	ggt Gly	2352
					tgc Cys 790											2400
					caa Gln											2448
					aca Thr											2496
					tgt Cys											2544
					gtg Val											2592

atg ttt gtt ttt ccg acc ctt tct tct gtc aag aaa tta gaa att tgg Met Phe Val Phe Pro Thr Leu Ser Ser Val Lys Lys Leu Glu Ile Trp	2640
865 870 875 880	
ggg gag gca gat gca aga ggt ttg agc tcc ata tct aat ctc agc act Gly Glu Ala Asp Ala Arg Gly Leu Ser Ser Ile Ser Asn Leu Ser Thr 885 890 895	2688
ctt aca tcc ctc aaa att ttc agt aac cac aca gtg act tca cta ctg Leu Thr Ser Leu Lys Ile Phe Ser Asn His Thr Val Thr Ser Leu Leu 900 905 910	2736
gaa gag atg ttc aaa agc ctc gaa aat ctc aaa tac ttg agt gtc tct Glu Glu Met Phe Lys Ser Leu Glu Asn Leu Lys Tyr Leu Ser Val Ser 915 920 925	2784
tac ttg gag aat ctc aaa gag ctg cct acc agc ctg gct agt ctc aat Tyr Leu Glu Asn Leu Lys Glu Leu Pro Thr Ser Leu Ala Ser Leu Asn 930 935 940	2832
aat ttg aag tgt ctg gat att cgt tat tgt tac gca cta gag agt ctc Asn Leu Lys Cys Leu Asp Ile Arg Tyr Cys Tyr Ala Leu Glu Ser Leu 945 950 955 960	2880
ccc gag gaa ggg ctg gaa ggt tta tct tca ctc aca gag tta ttt gtt Pro Glu Glu Gly Leu Glu Gly Leu Ser Ser Leu Thr Glu Leu Phe Val 965 970 975	2928
gaa cac tgt aac atg cta aaa tgt tta cca gag gga ttg cag cac cta Glu His Cys Asn Met Leu Lys Cys Leu Pro Glu Gly Leu Gln His Leu 980 985 990 .	2976
aca acc ctc aca agt tta aaa att cgg gga tgt cca caa ctg atc aag Thr Thr Leu Thr Ser Leu Lys Ile Arg Gly Cys Pro Gln Leu Ile Lys 995 1000 1005	3024
cgg tgt gag aag gga ata gga gaa gac tgg cac aaa att tct cac Arg Cys Glu Lys Gly Ile Gly Glu Asp Trp His Lys Ile Ser His 1010 1015 1020	3069
att cct aac gtg aat ata tat att taa Ile Pro Asn Val Asn Ile Tyr Ile 1025 1030	3096
<210> 20 <211> 1031 <212> PRT <213> Solanum bulbocastanum	
<400> 20	
Met Trp Leu Ser Thr Phe Pro Thr Ala Leu Ala Val Ala Val Ser Val 1 5 10 15	
Pro Ser Ser Leu Thr Pro Gln Asp Phe Ser Trp Arg Lys Phe Lys Leu 20 25 30	
Lys Arg Asn Val Lys Ile Ser Arg Asn Phe Glu Phe Ala Ile Ser Cys 35 40 45	
Ser Gly Asp Arg Ala Ala Ser Ile Gly Phe Asp Val Pro Phe Pro Lys 50 55 60	

Asp Tyr Thr Glu Leu Leu Gln Gln Val Phe Ile Leu Phe Ala Phe Ser Page 63

			WAR	F0235.ST	'25.txt	
65 <sup>-</sup>	70			75		80
Pro Leu Lys	Ile Gly Gly 85	Asp Gly	Glu Gly 90	Gly Ile	Glu Met Thi 95	Gly
Ser Ile Gln	Leu Ile Arg 100	Glu Phe	Cys Asp 105	Leu Leu	Val Ile Pro	Glu
Lys Ala Thr 115	Lys Thr Arg	Ile Phe 120	Phe Pro	Glu Ala	Asn Glu Val	Lys
Phe Ala Arg 130	Gln Ser Ile	Phe Gly 135	Gly Ala	Ser Phe 140	Lys Leu Asp	Tyr
Leu Thr Lys 145	Pro Ser Phe 150	Phe Glu	Asp Phe	Gly Phe 155	Thr Glu Lys	Val 160
Lys Met Ala	Asp Arg Val 165	Lys Pro	Glu Asp 170	Glu Leu	Phe Ile Val	
Tyr Pro Tyr	Phe Asn Val 180	Asn Gly	Glu Leu 185	Gly Leu	Val Phe Gly 190	/ Phe
Glu Lys Glu 195	Phe Lys Lys	Leu Ser 200	Ser Met	Phe Ser	Met Ile Glr 205	n Ala
Val Leu Glu 210	Asp Ala Gln	Glu Lys 215	Gln Leu	Lys Tyr 220	Lys Ala Ile	e Lys
Asn Trp Leu 225	GÎn Lys Leu 230	Asn Val	Ala Ala	Tyr Glu 235	Val Asp Asp	240
Leu Asp Asp	Cys Lys Thr 245	Glu Ala	Ala Arg 250	Phe Lys	Gln Ala Val 255	
Gly Arg Tyr	His Pro Arg 260	Thr Ile			Lys Val Gly 270	, Lys
Arg Met Lys 275	Glu Met Met	Glu Lys 280	Leu Asp	Ala Ile	Ala Glu Glu 285	ı Arg
Arg Asn Phe 290	His Leu Asp	Glu Arg 295	Ile Ile	Glu Arg 300	Gln Ala Ala	a Arg
Arg Gln Thr 305	Gly Phe Val 310	Leu Thr	Glu Pro	Lys Val 315	Tyr Gly Lys	320
Lys Glu Glu	Asp Glu Ile 325	Val Lys	Ile Leu 330	Ile Asn	Asn Val Ser 335	-

Ser Lys Glu Val Pro Val Leu Pro Ile Leu Gly Met Gly Gly Leu Gly 340 345 350

Lys	Thr	Thr 355	Leu	Ala	Gln	Met	Val 360	Phe	Asn	Asp	Gln	Arg 365	Ile	Thr	Glu
His	Phe 370	Asn	Leu	Lys	Ile	Trp 375	Val	Cys	Val	Ser	Asp 380	Asp	Phe	Asp	Glu
Lys 385	Arg	Leu	Ile	Lys	Ala 390	Ile	Val	Glu	Ser	Ile 395	Glu	Gly	Lys	Ser	Leu 400
Gly	Asp	Met	Asp	Leu 405	Ala	Pro	Leu	Gln	Lys 410	Lys	Leu	Gln	Glu	Leu 415	Leu
Asn	Gly	Lys	Arg 420	Tyr	Phe	Leu	Val	Leu 425	Asp	Asp	Val	Trp	Asn 430	Glu	Asp
Gln	Glu	Lys 435	Trp	Asp	Asn	Leu	Arg 440	Ala	Val	Leu	Lys	Ile 445	Gly	Ala	Ser
Gly	Ala 450	Ser	Ile	Leu	Ile	Thr 455	Thr	Arg	Leu	Glu	Lys 460	Ile	Gly	Ser	Ile
Met 465	Gly	Thr	Leu	Gln	Leu 470	Tyr	Gln	Leu	Ser	Asn 475	Leu	Ser	Gln	Glu	Asp 480
Cys	Trp	Leu	Leu	Phe 485	Lys	Gln	Arg	Ala	Phe 490	Cys	His	Gln	Thr	Glu 495	Thr
Ser	Pro	Lys	Leu 500	Met	Glu	Ile	Gly	Lys 505	Glu	Ile	Val	Lys	Lys 510	Cys	Gly
Gly	Val	Pro 515	Leu	Ala	Ala	Lys	Thr 520	Leu	Gly	Gly	Leu	Leu 525	Arg	Phe	Lys
Arg	Glu 530	Glu	Ser	Glu	Trp	Glu 535	His	Val	Arg	Asp	Ser 540	Glu	Ile	Trp	Asn
Leu 545	Pro	Gln	Asp	Glu	Asn 550	Ser	Val	Leu	Pro	Ala 555	Leu	Arg	Leu	Ser	Tyr 560
His	His	Leu	Pro	Leu 565	Asp	Leu	Arg	Gln	Cys 570	Phe	Ala	Tyr	Суѕ	Ala 575	Val
Phe	Pro	Lys	Asp 580	Thr	Lys	Ile	Glu	Lys 585	Glu	Tyr	Leu	Ile	Ala 590	Leu	Trp
Met	Ala	His 595	Ser	Phe	Leu	Leu	Ser 600	Lys	Gly	Asn	Met	Glu 605	Leu	Glu	Asp
Val	Gly 610	Asn	Glu	Val	Trp	Asn 615	Glu	Leu	Tyr	Leu	Arg 620	Ser	Phe	Phe	Gln
Glu	Ile	Glu	Val	Lys	Ser	Ala	Ser	Ala	Ser		Arg ge 6		Ile	Arg	Gln

Ile	Asn	Val	Lys	Asp 645	Asp	Glu	Asp	Met	Met 650	Phe	Ile	Val	Thr	Asn 655	Tyr
Lys	Asp	Met	Met 660	Ser	Ile	Gly	Ser	Pro 665	Lys	Trp	Cys	Leu	Leu 670	Thr	Leu
Leu	Arg	Ser 675	Leu	Lys	Gly	Glu	Arg 680	Lys	Gly	Tyr	Gln	Leu 685	Gly	Glu	Leu
Arg	Asn 690	Leu	Asn	Leu	Arg	Gly 695	Ala	Ile	Ser	Ile	Thr 700	His	Leu	Glu	Arg
Val 705	Lys	Asn	Asp	Met	Glu 710	Ala	Lys	Glu	Ala	Asn 715	Leu	Ser	Ala	Lys	Ala 720
Asn	Leu	His	Ser	Leu 725	Ser	Met	Ser	Trp	Asp 730	Arg	Pro	Asn	Arg	Tyr 735	Glu
Ser	Glu	Glu	Val 740	Lys	Val	Leu	Glu	Ala 745	Leu	Lys	Pro	His	Pro 750	Asn	Leu
Lys	Tyr	Leu 755	Glu	Ile	Ile	Asp	Phe 760	Cys	Gly	Phe	Cys	Leu 765	Pro	Asp	Trp
Met	Asn 770	His	Ser	Val	Leu	Lys 775	Asn	Val	Val	Ser	Ile 780	Leu	Ile	Ser	Gly
Cys 785	Glu	Asn	Cys	Ser	Cys 790	Leu	Pro	Pro	Phe	Gly 795	Glu	Leu	Pro	Cys	Leu 800
Glu	Ser	Leu	Glu	Leu 805	Gln	Asp	Gly	Ser	Val 810	Glu	Val	Glu	Phe	Val 815	Glu
Asp	Ser	Gly	Phe 820	Pro	Thr	Arg	Arg	Arg 825	Phe	Pro	Ser	Leu	Arg 830	Lys	Leu
His	Ile	Gly 835	Gly	Phe	Cys	Asn	Leu 840	Lys	Gly	Leu	Gln	Arg 845	Met	Glu	Gly
Glu	Glu 850	Gln	Phe	Pro	Val	Leu 855	Glu	Glu	Met	Lys	Ile 860	Ser	Asp	Cys	Pro
Met 865	Phe	Val	Phe	Pro	Thr 870	Leu	Ser	Ser	Val	Lys 875	Lys	Leu	Glu	Ile	Trp 880
Gly	Glu	Ala	Asp	Ala 885	Arg	Gly	Leu	Ser	Ser 890	Ile	Ser	Asn	Leu	Ser 895	Thr
Leu	Thr	Ser	Leu 900	Lys	Ile	Phe	Ser	Asn 905	His	Thr	Val	Thr	Ser 910	Leu	Leu

Glu	Glu	Met 915	Phe	Lys	Ser	Leu	Glu 920	Asn	Leu	Lys	Tyr	Leu 925	Ser	Val	Ser	
Tyr	Leu 930	Glu	Asn	Leu	Lys	Glu 935	Leu	Pro	Thr	Ser	Leu 940	Ala	Ser	Leu	Asn	
Asn 945	Leu	Lys	Cys	Leu	Asp 950	Ile	Arg	Tyr	Суѕ	Tyr 955	Ala	Leu	Glu	Ser	Leu 960	
Pro	Glu	Glu	Gly	Leu 965	Glu	Gly	Leu	Ser	Ser 970	Leu	Thr	Glu	Leu	Phe 975	Val	
Glu	His	Cys	Asn 980	Met	Leu	Lys	Cys	Leu 985	Pro	Glu	Gly	Leu	Gln 990	His	Leu	
Thr	Thr	Leu 995	Thr	Ser	Leu	Lys	Ile 1000		g Gly	у Су:	s Pro	Gl: 100		eu II	le Lys	
Arg	Cys 1010		ı Lys	s Gly	/ Ile	e Gly 101		lu As	sp Ti	rp Hi	is Ly 10	/s : )20	Ile S	Ser H	His	
Ile	Pro 1025		n Val	l Ası	n Ile	€ Tyr 103		Le								
<210 <211 <212 <213	L> 2 2> I	21 2967 DNA Solar	num k	oulbo	ocast	canum	n									
<220 <221 <222	L> (	CDS (1)	. (296	57)												
<400 atg Met 1	gcg	21 gaa Glu	gct Ala	ttt Phe 5	ctt Leu	caa Gln	gtt Val	ctg Leu	cta Leu 10	gaa Glu	aat Asn	ctc Leu	act Thr	tct Ser 15	ttc Phe	48
		Asp	Lys	Leu	Val	Leu	Ile	Phe	Gly	Phe	gaa Glu	Lys	Glu	Cys		96
											gtg Val					144
											aat Asn 60					192
											ttg Leu					240
											Gly ggg					288
											agg Arg					336

Page 67

atg gag aaa cta gat gca att gct gag gaa agg aag ttt cat ttc 384 Met Glu Lys Leu Asp Ala Ile Ala Glu Glu Arg Arg Lys Phe His Phe 120 ctt gaa aaa att acg gag aga caa gct gcc gct gct acg cgt qaa aca 432 Leu Glu Lys Ile Thr Glu Arg Gln Ala Ala Ala Ala Thr Arg Glu Thr 135 ggt ttt gtg tta act gaa cca aaa gtc tac gga agg gac aaa gag gag 480 Gly Phe Val Leu Thr Glu Pro Lys Val Tyr Gly Arg Asp Lys Glu Glu 150 gat gag ata gtg aaa att ctg ata aac aat gtt aat gtt gcc gaa gaa 528 Asp Glu Ile Val Lys Ile Leu Ile Asn Asn Val Asn Val Ala Glu Glu ctt cca gtc ttc cct ata att ggt atg ggg gga cta gga aag acg aca 576 Leu Pro Val Phe Pro Ile Ile Gly Met Gly Gly Leu Gly Lys Thr Thr 185 ctt gcc caa atg atc ttc aac gat gag aga gta act aag cat ttc aat 624 Leu Ala Gln Met Ile Phe Asn Asp Glu Arg Val Thr Lys His Phe Asn 200 ccc aaa ata tgg gtt tgt gtc tca gat gat ttt gat gag aag agg tta 672 Pro Lys Ile Trp Val Cys Val Ser Asp Asp Phe Asp Glu Lys Arg Leu att aag aca att ata gga aat att gaa aga agt tot oot cat gtt gag 720 Ile Lys Thr Ile Ile Gly Asn Ile Glu Arg Ser Ser Pro His Val Glu 230 240 gac ttg gct tca ttt cag aag aag ctc cag gag tta ttg aat gga aaa 768 Asp Leu Ala Ser Phe Gln Lys Lys Leu Gln Glu Leu Leu Asn Gly Lys cga tac ttg ctt gtc tta gat gat gtt tgg aat gat gat cta gaa aag Arg Tyr Leu Leu Val Leu Asp Asp Val Trp Asn Asp Asp Leu Glu Lys 816 tgg gct aag tta aga gca gtc tta act gtt gga gca aga ggt gct tct Trp Ala Lys Leu Arg Ala Val Leu Thr Val Gly Ala Arg Gly Ala Ser 864 275 280 att cta gct act cgt ctt gaa aag gtt gga tca att atg gga acg 912 Ile Leu Ala Thr Thr Arg Leu Glu Lys Val Gly Ser Ile Met Gly Thr tcg caa cca tat cat ttg tca aat ttg tct cca cat gat agt tta ctt 960 Ser Gln Pro Tyr His Leu Ser Asn Leu Ser Pro His Asp Ser Leu Leu 310 315 ttg ttt atg caa cgc gca ttt ggg caa caa aaa gaa gca aat cct aat 1008 Leu Phe Met Gln Arg Ala Phe Gly Gln Gln Lys Glu Ala Asn Pro Asn 335 1056 tta gcg gcc aag act ctt ggt ggt ctt tta cgc ttc aag aga gaa gag 1104 Leu Ala Ala Lys Thr Leu Gly Gly Leu Leu Arg Phe Lys Arg Glu Glu 360 agt gaa tgg gaa cat gtg aga gat aat gag att tgg agt ctg cct caa 1152 Ser Glu Trp Glu His Val Arg Asp Asn Glu Ile Trp Ser Leu Pro Gln

											5.ST					
gat Asp 385	gaa Glu	agt Ser	tct Ser	att Ile	ttg Leu 390	cct Pro	gct Ala	cta Leu	aga Arg	ctg Leu 395	agt Ser	tat Tyr	cat His	cac His	ctt Leu 400	1200
	ctt Leu															1248
	acc Thr															1296
	ttt Phe															1344
_	gta Val 450			_			_						-		_	1392
	aaa Lys															1440
	gct Ala															1488
	ata Ile															1536
	gtg Val							_		_		_		_	_	1584
	agg Arg 530															1632
	att Ile															1680
	ttc Phe	_	_					_	_	_					_	1728
	ctt Leu	_	_			_					-	_				1776
	agt Ser			_	_		-			-	_	_		_		1824
	act Thr 610								_	_		_		_		1872
	ggt Gly						-	_							-	1920
	aaa Lys															1968
	gtg Val															2016

								•								
				tta Leu												2064
				gaa Glu												2112
				tta Leu												2160
				cac His 725												2208
				aac Asn												2256
				cta Leu												2304
				gat Asp												2352
		_		aaa Lys		_					_	_	_			2400
				gaa Glu 805												2448
				tgc Cys		_		-							_	2496
_			-	gtt Val									_	_		2544
				agc Ser												2592
	_	_		ctc Leu		_					-					2640
-		_	_	ttc Phe 885		_		_				_	_			2688
_	_		_	ctc Leu			_	_	-				_	_	-	2736
-				agt Ser			-				-					2784
			_	ttt Phe	-			_	_	_			_			2832

									WAR	F023	5.ST	25. t	xt				
				cac His					aca	aat	tta	gga	gtt			288	0
				gaa Glu 965												292	8
				cac His								tag				296	7
<210 <211 <212 <213	L>	22 988 PRT Solar	num l	oulbo	ocast	anur	n										
<400	)> 2	22															
Met 1	Ala	Glu	Ala	Phe 5	Leu	Gln	Val	Leu	Leu 10	Glu	Asn	Leu	Thr	Ser 15	Phe		
Ile	Gly	Asp	Lys 20	Leu	Val	Leu	Ile	Phe 25	Gly	Phe	Glu	Lys	Glu 30	Cys	Glu		
Lys	Leu	Ser 35	Ser	Val	Phe	Ser	Thr 40	Ile	Gln	Ala	Val	Val 45	Gln	Asp	Ala		
Gln	Glu 50	Lys	Gln	Leu	Lys	Asp 55	Lys	Ala	Ile	Glu	Asn 60	Trp	Leu	Gln	Lys		
Leu 65	Asn	Ser	Ala	Ala	Tyr 70	Glu	Val	Asp	Asp	Ile 75	Leu	Gly	Glu	Cys	Lys 80		
Asn	Glu	Ala	Ile	Arg 85	Phe	Glu	Gln	Ser	Arg 90	Leu	Gly	Phe	Tyr	His 95	Pro		
Gly	Ile	Ile	Asn 100	Phe	Arg	His	Lys	Ile 105	Gly	Arg	Arg	Met	Lys 110	Glu	Ile		
Met	Glu	Lys 115	Leu	Asp	Ala	Ile	Ala 120	Glu	Glu	Arg	Arg	Lys 125	Phe	His	Phe		
Leu	Glu 130	Lys	Ile	Thr	Glu	Arg 135	Gln	Ala	Ala	Ala	Ala 140	Thr	Arg	Glu	Thr		
Gly 145	Phe	Val	Leu	Thr	Glu 150	Pro	Lys	Val	Tyr	Gly 155	Arg	Asp	Lys	Glu	Glu 160		

Asp Glu Ile Val Lys Ile Leu Ile Asn Asn Val Asn Val Ala Glu Glu

165

Pro	Lys 210	Ile	Trp	Val	Cys	Val 215	Ser	Asp	Asp	Phe	Asp 220	Glu	Lys	Arg	Leu
Ile 225	Lys	Thr	Ile	Ile	Gly 230	Asn	Ile	Glu	Arg	Ser 235	Ser	Pro	His	Val	Glu 240
Asp	Leu	Ala	Ser	Phe 245	Gln	Lys	Lys	Leu	Gln 250	Glu	Leu	Leu	Asn	Gly 255	Lys
Arg	Tyr	Leu	Leu 260	Val	Leu	Asp	Asp	Val 265	Trp	Asn	Asp	Asp	Leu 270	Glu	Lys
Trp	Ala	Lys 275	Leu	Arg	Ala	Val	Leu 280	Thr	Val	Gly	Ala	Arg 285	Gly	Ala	Ser
Ile	Leu 290	Ala	Thr	Thr	Arg	Leu 295	Glu	Lys	Val	Gly	Ser 300	Ile	Met	Gly	Thr
Ser 305	Gln	Pro	Tyr	His	Leu 310	Ser	Asn	Leu	Ser	Pro 315	His	Asp	Ser	Leu	Leu 320
Leu	Phe	Met	Gln	Arg 325	Ala	Phe	Gly	Gln	Gln 330	Lys	Glu	Ala	Asn	Pro 335	Asn
Leu	Val	Ala	Ile 340	Gly	Lys	Glu	Ile	Val 345	Lys	Lys	Cys	Gly	Gly 350	Val	Pro
Leu	Ala	Ala 355	Lys	Thr	Leu	Gly	Gly 360	Leu	Leu	Arg	Phe	Lys 365	Arg	Glu	Glu
Ser	Glu 370	Trp	Glu	His	Val	Arg 375	Asp	Asn	Glu	Ile	Trp 380	Ser	Leu	Pro	Gln
Asp 385	Glu	Ser	Ser	Ile	Leu 390	Pro	Ala	Leu	Arg	Leu 395	Ser	Tyr	His	His	Leu 400
Pro	Leu	Asp	Leu	Arg 405	Gln	Cys	Phe	Ala	Tyr 410	Cys	Ala	Val	Phe	Pro 415	Lys
Asp	Thr	Lys	Met 420	Ile	Lys	Glu	Asn	Leu 425	Ile	Thr	Leu	Trp	Met 430	Ala	His
Gly	Phe	Leu 435	Leu	Ser	Lys	Gly	Asn 440	Leu	Glu	Leu	Glu	Asp 445	Val	Gly	Asn
Glu	Val 450	Trp	Asn	Glu	Leu	Tyr 455	Leu	Arg	Ser	Phe	Phe 460	Gln	Glu	Ile	Glu
Ala 465	Lys	Ser	Gly	Asn	Thr 470	Tyr	Phe	Lys	Ile	His 475	Asp	Leu	Ile	His	Asp 480

Leu	Ala	Thr	Ser	Leu 485	Phe	Ser	Ala	Ser	Ala 490	Ser	Суѕ	Gly	Asn	Ile 495	Arg

Glu Ile Asn Val Lys Asp Tyr Lys His Thr Val Ser Ile Gly Phe Ser 500 505 510

Ala Val Val Ser Ser Tyr Ser Pro Ser Leu Leu Lys Lys Phe Val Ser 515 520 525

Leu Arg Val Leu Asn Leu Ser Tyr Ser Lys Leu Glu Gln Leu Pro Ser 530 540

Ser Ile Gly Asp Leu Leu His Leu Arg Tyr Leu Asp Leu Ser Cys Asn 545 550 555 560

Asn Phe Arg Ser Leu Pro Glu Arg Leu Cys Lys Leu Gln Asn Leu Gln 565 570 575

Thr Leu Asp Val His Asn Cys Tyr Ser Leu Asn Cys Leu Pro Lys Gln
580 585 590

Thr Ser Lys Leu Ser Ser Leu Arg His Leu Val Val Asp Gly Cys Pro 595 600 605

Leu Thr Ser Thr Pro Pro Arg Ile Gly Leu Leu Thr Cys Leu Lys Thr 610 615 620

Leu Gly Phe Phe Ile Val Gly Ser Lys Lys Gly Tyr Gln Leu Gly Glu 625 630 635

Leu Lys Asn Leu Asn Leu Cys Gly Ser Ile Ser Ile Thr His Leu Glu 645 650 655

Arg Val Lys Asn Asp Thr Asp Ala Glu Ala Asn Leu Ser Ala Lys Ala 660 665 670

Asn Leu Gln Ser Leu Ser Met Ser Trp Asp Asn Asp Gly Pro Asn Arg 675 680 685

Tyr Glu Ser Glu Glu Val Lys Val Leu Glu Ala Leu Lys Pro His Pro 690 695 700

Asn Leu Lys Tyr Leu Glu Ile Ile Ala Phe Gly Gly Phe Arg Phe Pro
705 710 715 720

Ser Trp Ile Asn His Ser Val Leu Glu Lys Val Ile Ser Val Arg Ile 725 730 735

Lys Ser Cys Lys Asn Cys Leu Cys Leu Pro Pro Phe Gly Glu Leu Pro

Cys Leu Glu Asn Leu Glu Leu Gln Asn Gly Ser Ala Glu Val Glu Tyr
755 760 765

vai	770	GIU	Asp	ASP	vai	775	ser	Arg	rne	ser	780	Arg	Arg	Ser	Pne		
Pro 785	Ser	Leu	Lys	Lys	Leu 790	Arg	Ile	Trp	Phe	Phe 795	Arg	Ser	Leu	Lys	Gly 800		
Leu	Met	Lys	Glu	Glu 805	Gly	Glu	Glu	Lys	Phe 810	Pro	Met	Leu	Glu	Glu 815	Met		
Ala	Ile	Leu	Tyr 820	Cys	Pro	Leu	Phe	Val 825	Phe	Pro	Thr	Leu	Ser 830	Ser	Val		
Lys	Lys	Leu 835	Glu	Val	His	Gly	Asn 840	Thr	Asn	Thr	Arg	Gly 845	Leu	Ser	Ser		
Ile	Ser 850	Asn	Leu	Ser	Thr	Leu 855	Thr	Ser	Leu	Arg	Ile 860	Gly	Ala	Asn	Tyr		
Arg 865	Ala	Thr	Ser	Leu	Pro 870	Glu	Glu	Met	Phe	Thr 875	Ser	Leu	Thr	Asn	Leu 880		
Glu	Phe	Leu	Ser	Phe 885	Phe	Asp	Phe	Lys	Asn 890	Leu	Lys	Asp	Leu	Pro 895	Thr		
Ser	Leu	Thr	Ser 900	Leu	Asn	Ala	Leu	Lys 905	Arg	Leu	Gln	Ile	Glu 910	Ser	Cys		
Asp	Ser	Leu 915	Glu	Ser	Phe	Pro	Glu 920	Gln	Gly	Leu	Glu	Gly 925	Leu	Thr	Ser		
Leu	Thr 930	Gln	Leu	Phe	Val	Lys 935	Tyr	Cys	Lys	Met	Leu 940	Lys	Cys	Leu	Pro		
Glu 945	Gly	Leu	Gln	His	Leu 950	Thr	Ala	Leu	Thr	Asn 955	Leu	Gly	Val	Ser	Gly 960		
Cys	Pro	Glu	Val	Glu 965	Lys	Arg	Суѕ	Asp	Lys 970	Glu	Ile	Gly	Glu	Asp 975	Trp		
His	Lys	Ile	Ala 980	His	Ile	Pro	Asn	Leu 985	Asp	Ile	His						
<21 <21 <21 <21	1> 2 2> [	23 2000 ONA Artif	ficia	al Se	equer	nce											
<22 <22		Promo	oter														
<400 acg		23 ctt a	aataa	aaaa	ag aa	ataat	caaaa	a tta	atato	gata	attt	ttat	caa t	tacaa	atggcc		60
ttt	atato	gat ç	gaaaa	aaaa	aa ga	aaga	aatt	aga	atgad				aaa a	aataa	atctta	1.	20
										ra	ge 7	4					

aàgaattacg	atttatatat	aataaaatta	aatttaaaat	ttgatgaaaa	aatagagaaa	180
agaggaagat	gatgaagtga	aatgacgtgg	tggtgggtcc	atgtgacata	aaaaaaatt	240
ctcttaaata	atcctttcat	actaatgata	aaatttttt	tttttttt	tttttactaa	300
ttgcgtatag	agaaaaggaa	aatggggcgg	taattacaaa	gtagggaatc	gaactttatc	360
aacaagttga	gagttcaagt	aatcaaccaa	ctaaactact	aaaattttc	taattaatga	420
taattgtaat	tcatttagca	taaaaaattt	cattgcactt	acttttagag	ttttgaaaac	480
agtacttcat	ctattctata	ttaattaaat	tttctatatt	aattaaattt	gtgaggtaat	540
acaaacttat	taagaaaaat	atttaaggac	ataatttaac	tcatatttt	cactattgtt	600
ttttgtgaaa	tcataaatat	aactttgtaa	atagtgcaat	ttatctccta	gaagcaaatt	660
tcaccaaaga	aaagggcaaa	gatggaaaag	aaactaaata	ttcatcttaa	actttgaaca	720
attcaattat	tttgaacaat	gaaaaaaatc	tcaaaaattc	aattaatatg	aaatggagag	780
agtaacttta	ttttagaggc	aaaaaattag	tactccatcc	gttcactttt	atttgtcatg	840
ttgcgctttt	cgaaagtcaa	tttgactaat	ttttaaagct	aaattagatt	acactaattc	900
aatatttaa	acagaaaaat	tagatattca	aaaactatac	aaaaaatatt	atacattgca	960
attttttgca	tatcaatatg	ataaaaaaat	atattgtaaa	atattagtca	aaattttat	1020
agtttgactc	taatcatgaa	aagtataata	attaatagtg	gacggaggaa	gtattgtctt	1080
tccagatttg	tggccatttt	tgggccaagg	gccattagca	gttctcttca	ttttctactt	1140
ctgtctcata	ttagatgggc	atcttactaa	aaatatttgt	ctcatattac	ttgattattt	1200
attaaatcaa	aaagaattaa	ttaattttt	ctcattttac	ccctacaatt	aatatagttt	1260
taaaagtttt	aaacaaattt	tgaagaatca	aaatttcttt	ttgcaagaga	cttattaata	1320
taaacaaagg	ataaaataat	aaaatttgtc	aatttattga	cgatcactta	ataatcatat	1380
aaaatagaat	atgtttatct	aatatgagac	ggagaaaata	tatcctaaaa	tatttttgga	1440
cagatatgtg	atattctaac	cattcactag	actatattat	gcattttagc	cgccaatgac	1500
ttatttcagc	tttaattaat	taggaaagag	gaaactgcca	atgaggaaga	gtaggggcgt	1560
agttgctgtc	gacgaaaaaa	agataatact	cactcttttc	gatttttatt	tttatttatc	1620
acttttaacc	tatcatgtaa	aaagataatt	attttttca	tgctttatcc	ttagtattaa	1680
acaatttaat	agggattatt	ttgtaaaata	tttatatgaa	taattgtttt	cgtaatgaat	1740
ttgtccggtc	aaacaatgat	aaataaaaat	gaatgaagag	agtagaaaac	aaaacaaaag	1800
aacaagttga	caacttgaga	gattaaaagg	gtccaaaacg	ccttggattt	tgagattcca	1860
tatgtgaaat	ttccatgaaa	taattgaatt	tgtattatta	caagtcaaac	tttccatttc	1920
attccaacta	gccatcttgg	tttcaaaatt	acacattcat	tcattcacag	atctaatatt	1980
cttaatagtg	atttccacat					2000

<sup>&</sup>lt;210> 24 <211> 32 <212> DNA <213> Artificial Sequence

<220> <223>	Primer	
<400> ctacct	24 tgta attaccgccc cattttcctt tt	32
<210><211><211><212><213>	25 32 DNA Artificial Sequence	
<220> <223>	Primer	
<400> tgtcac	25 ataa attgacacaa agggagtact tg	32
<210><211><211><212><213>	26 32 DNA Artificial Sequence	
<220> <223>	Primer	
<400> aagcgg	26 aagg aatggtttgg gtatgataaa at	32
<210><211><211><212><213>	27 32 DNA Artificial Sequence	
<220> <223>	Primer	
<400> ctttca	27 aaat gggaaaggag attaaaggtg ga	32
<210><211><211><212><213>	28 33 DNA Artificial Sequence	
<220> <223>	Primer	
<400> ccttgt	28 aatt gctaccccat tagtatgaaa gga	33
<210><211><211><212><212><213>	29 32 DNA Artificial Sequence	
<220> <223>	Primer	
<400> aggaga	29 ttaa aggtggataa tcaaacctca cc	32
<210> <211>	30 32	

<212>	DNA	
<213>	Artificial Sequence	
<220> <223>	Primer	
<400>	30	
gtttctt	tttc catctttgcc cttttcttta gg	32
<210> <211>	31 32	
	DNA	
<213>	Artificial Sequence	
<220>		
<223>	Primer	
<400>	31	
atcaatt	ttta tcatacccaa accattcctt cc	32
<210> <211>	32 35	
	DNA	
<213>	Artificial Sequence	
<220>		
<223>	Primer	
<400>	32	
ttgactt	tgag tacaacaaat ccaagtataa agaag	35
10101		
<210> <211>	33 32	
<212>	DNA	
<213>	Artificial Sequence	
<220> <223>	Drimor	
\2237	Primer	
<400>	33 gcaa gtttagaaaa tggtcatgat ac	32
geguage	geau geeeugaaau eggeeutgae ac	32
<210>	34	
<211>	34	
<212> <213>	DNA Artificial Sequence	
<220> <223>	Primer	
<400> tgattat	34 taac agacactcaa ttagcaagcc tgtg	34
-	-	
<210>	35	
<211>	32 DNA	
<212> <213>	DNA Artificial Sequence	
<220>	-	
<223>	Primer	
<400>	35	
	aaaa qaactcaaag cggaaggaat gg	32

1

<210>	36	
<211>	10	
<212>	DNA	
<213>	Artificial Sequence	
<220>		
<223>	Primer	
	•	
<400>	36	
ggcact	gagg	10
<210>	37	
<211>	25	
<212>	DNA	
<213>	Artificial Sequence	
<220>		
<223>	Primer	
12237	TITMOI	
<400>	37	
		25
CCCLLC	ttat tctttaagca aactt	25
4010s	30	
<210>	38	
<211>	19	
<212>	DNA	
<213>	Artificial Sequence	
<220>		
<223>	Primer	
<400>	38	
attttt	ccc tcccagctt	19
	· •	
<210>	39	
<211>	20	
<212>	DNA	
<213>	Artificial Sequence	
\213/	Altilitial Sequence	
<220N		
<220>	D. dans.	
<223>	Primer	
<400>	39	
attttg	ctcg aatcgctcat ·	20
<210>	40	
<211>	21	
<212>	DNA	
<213>	Artificial Sequence	
	•	
<220>		
<223>	Primer	
<400>	40	
	tgg atcacttgtc t	21
	55	ھ عد
<210>	41	
<211>	21	
<211>		
	DNA	
<213>	Artificial Sequence	
Z2205		
<220>	Desire	
<223>	Primer	
<1000	4.1	
<400>	41	
tgtttgi	gca ttgaagattg g	21

<210> <211> <212> <213>		
<220> <223>	Primer	
<400> tggtaa	42 gaag ggcattccat a	21
<211> <212>		
<220> <223>	Primer	
<400> ggaaat	43 acta gaggggaggg agt	23
<210><211><211><212><213>		
<220> <223>	Primer	
<400> tgaata	44 agca gttcggtttg aa	22
<210><211><211><212><213>	45 20 DNA Artificial Sequence	
<220> <223>	Primer	
<400> tctcttc	45 ggga tcacgattca	20
<210><211><211><212><213>	46 20 DNA Artificial Sequence	
<220> <223>	Primer	
<400> tttaat	46 ttcg gcggatgaac	20
<210><211><211><212><212><213>	47 23 DNA Artificial Sequence	
<220> <223>	Primer	

<400> tttgga	47 ggat agcaatactt gga	23
<210><211><211><212><213>	48 23 DNA Artificial Sequence	
<220> <223>	Primer	
<400> agcaac	48 tggt gagaaaatgt ctt	23
<210> <211> <212> <213>	49 20 DNA Artificial Sequence	
<220> <223>	Primer	
<400> cggttc	49 agct gacctttcat	20
<210> <211> <212> <213>	50 20 DNA Artificial Sequence	
<220> <223>	Primer	
<400> acctgc	50 gagt ggatcaaaac	20
<210> <211> <212> <213>	51 24 DNA Artificial Sequence	
<220> <223>	Primer	
<400> tcacaa	51 tgct aatatgtggt ttga	24
<210><211><211><212><213>	52 20 DNA Artificial Sequence	
<220> <223>	Primer	
<400> agttgt	52 ctgt ggctgccatt	20
<210> <211> <212> <213>	53 21 DNA Artificial Sequence	

<223>	Primer	
<400>	53 ccaa gtgaaaagtc g	21
<210><211><211><212><212><213>	19 DNA	
<220> <223>	Primer	
<400> atcaag	54 cacc tccccaaac	19
<210><211><211><212><213>	55 20 DNA Artificial Sequence	
<220> <223>	Primer	
<400> aactag	55 cccg cgatcaacta	20
<210><211><211><212><213>	56 20 DNA Artificial Sequence	
<220> <223>	Primer	
<400> aaaccg	56 acac agatgcaaca	20
<210><211><211><212><213>	57 20 DNA Artificial Sequence	
<220> <223>	Primer	
<400> ccctct	57 gttc cgtgacaaat	20
<210><211><211><212><213>	58 20 DNA Artificial Sequence	
<220> <223>	Primer	
<400> cacagaa	58 aggg ggttgatctc	20
<210><211><211><212><213>	59 26 DNA Artificial Sequence	

<220> <223>	Primer	
<400> tgagtt	59 ccac agtctgtaca taacaa	26
<210><211><211><212><213>	60 22 DNA Artificial Sequence	
<220> <223>	Primer	
<400> tttctt	60 cete tecettetee tt	22
<210> <211> <212> <213>	61 20 DNA Artificial Sequence	
<220> <223>	Primer	
<400> aacaaga	61 atga gootggtgtg	20
<210><211><211><212><213>	62 20 DNA Artificial Sequence	
<220> <223>	Primer	
<400> atcaca	62 cccc agaggcaaaa	20
<210><211><211><212><213>	63 20 DNA Artificial Sequence	
<220> <223>	Primer	
<400> atcaate	63 ccat catgtgagca	20
<210><211><212><212><213>	64 22 DNA Artificial Sequence	
<220> <223>	Primer	
<400> tcagaaa	64 aata agcacgttga ca	22
<210> <211>	65 20	

10101	D.1.7	WART 0233.3123. CAL	
<212> <213>	DNA Artificial Sequence		
<220>	•		
<223>	Primer		
<400>	65		0.0
cttgag	aagg caacgacaga		20
<210>	66		
<211>	20		
<212> <213>	DNA Artificial Sequence		
<220>			
<223>	Primer		
	66		
gaaggc	gggt aaacagacag		20
<210>	67		
<211>	20		
<212> <213>	DNA Artificial Sequence		
<220>	•		
<223>	Primer		
<400>	67		
caatcg	ctcc ttccaacttc		20
.010.	60		
<210> <211>	68 20		
<212> <213>	DNA Artificial Seguence		
	Artificial Sequence		
<220> <223>	Primer		
<400>	68		
	gcat tcgaagaaaa		20
<210> <211>	69 24		
<212>	DNA		
	Artificial Sequence		
<220> <223>	Primer		
<400>	69		
	ctcc tcaagttcta caca		24
<210>	70		
<211> <212>	19 DNA		
<213>	Artificial Sequence		
<220>			
<223>	Primer		
<400> gatacg	70 ggtg ccaggattc		19

<210>	71	
<211>	20	
<212>	DNA	_
<213>	Artificial	Sequence
<220>		
<223>	Primer	
12237	TITMOI	
<400>	71	
	cagc aagttca	agc
		-
<210>	72	
<211>	20	
<212>	DNA	
<213>	Artificial	Sequence
<220>		
<223>	Primer	
	7.0	
<400>	72	
ttattg	tcca tgtcgct	cca
Z2105	72	
<210> <211>	73	
<211>	20 DNA	
<212>	DNA Artificial	Seguence
<b>\L13/</b>	Artificial	sequence
<220>		
<223>	Primer	
	LLIMOL	
<400>	73	
	atgc ctaaaco	caga
50000		
<210>	74	
<211>	20	
<212>	DNA	
<213>	Artificial	Sequence
		- 4
<220>		
<223>	Primer	
<400>	74	
atcgcc	cgct caactta	ata
=		
<210>	75	
<211>	20	
<212>	DNA	
<213>	Artificial	Sequence
<220>		
<223>	Primer	
1400	7.5	
<400>	75	
tgaggt.	attg ctgtggg	ıttg
Z21.05	7.0	
<210>	76	
<211>	21	-
<212>	DNA	<b>Q</b>
<213>	Artificial	Sequence
Z200:		
<220>	David manage	
<223>	Primer	
Z4005	76	
<400>	76	rtan n
Luaatt	cage ceagaac	jiga a

<210> <211> <212> <213>	77 20 DNA Artificial Sequence	
<220> <223>	Primer	
	77 tcgc ctgtccaaac	20
<210><211><211><212><213>		
<220> <223>	Primer	
	78 tctc gttggatgaa	20
<210><211><211><212><213>		
<220> <223>	Primer	
<400> ccctcg	79 acat gaaccagaag	20
<210><211><211><212><213>	80 20 DNA Artificial Sequence	
<220> <223>	Primer	
<400> tgtcca	80 tgta ggccaagacc	20
<210><211><211><212><213>	81 20 DNA Artificial Sequence	
<220> <223>	Primer	
<400> acaggc	81 cagg gttcaaaata	20
<210><211><211><212><213>	82 20 DNA Artificial Sequence	
<220> <223>	Primer	

<400> gcaatg	82 gaca gacttgatgc	20
<210><211><211><212><213>	83 21 DNA Artificial Sequence	
<220> <223>	Primer	
<400> gaggca	83 aacc ctctgttccg t	21
<210><211><211><212><213>	84 21 DNA Artificial Sequence	
<220> <223>	Primer	
<400> gctcca	84 agtg gaggaaatgc c	21
<210><211><211><212><213>	85 20 DNA Artificial Sequence	
<220> <223>	Primer	
<400> aacaag	85 atga gcctggtgtg	20
<210><211><212><212><213>	86 20 DNA Artificial Sequence	
<220> <223>	Primer	
<400> atcaca	86 tccc agaggcaaaa	20
<210><211><211><212><213>	87 18 DNA Artificial Sequence	
<220> <223>	Primer	
<400> gcttag	87 tgcc cttaagcg	18
<210> <211> <212> <213>	88 18 DNA Artificial Sequence	
<220>		

<223>	Primer	
<400> ctgact	88 aacc ggatggcc	18
<210><211><211><212><213>	89 26 DNA Artificial Sequence	
<220> <223>	Primer	
<400> aaaatt	89 gtcc tcctctaatt ttcttt	26
<210> <211> <212> <213>	90 23 DNA Artificial Sequence	
<220> <223>	Primer	
<400> tgatat	90 gaaa agaaagtggt tgc	23
<210> <211> <212> <213>	91 21 DNA Artificial Sequence	
<220> <223>	Primer	
<400> cgtgaa	91 gtga aatgctcaac a	21
<210> <211> <212> <213>	92 20 DNA Artificial Sequence	
<220> <223>	Primer .	
<400> gcaaac	92 tttg gaaggattcg	20
<210> <211> <212> <213>	93 20 DNA Artificial Sequence	
<220> <223>	Primer	
<400> cctgag	93 cctc ggtgagagta	20
<210><211><211><212><212><213>	94 20 DNA Artificial Sequence	

<220> <223>	Primer	
<400> acccaa	94 aact cccaacctct	20
<210><211><211><212><213>	DNA	
<220> <223>	Primer	
<400> ctggtt	95 tgac aatgctggtg	20
<210><211><211><212><213>	DNA	
<220> <223>	Primer	
<400> gacacte	96 caag gctgccattt	20
<210><211><212><212><213>		
<220> <223>	Primer	
<400> tctgca	97 gaaa accatctcag g	21
<210><211><211><212><213>	98 20 DNA Artificial Sequence	
<220> <223>	Primer	
<400> agctct	98 taac acgcctggaa	20
<210><211><211><212><213>	99 20 DNA Artificial Sequence	
<220> <223>	Primer	
<400> gcttaa	99 gcac gcttctgaca	20
<210> <211>	100 20	

		WARE 0233.3123.CXC	
<212> <213>	DNA Artificial Sequence		
<220> <223>	Primer		
<400>	100 gacc agccattgat		20
tgacat	gace agecactgat		20
<210> <211>	101 20		
<212> <213>	DNA Artificial Sequence		
<220> <223>	Primer		
<400>	101 aaaa gaacccatct		20
			20
<210> <211>	102 20		
<212> <213>	DNA Artificial Sequence		
<220> <223>	Primer		
<400>	102 acct ctgtcttctc		20
3 3	j		
<210> <211>	103 20		
<212> <213>	DNA Artificial Sequence		
<220> <223>	Primer		
<400>	103 ggcc attgaaagaa		20
<210> <211>	104 20		
<212> <213>	DNA Artificial Sequence		
<220> <223>	Primer	:	
<400> ttggate	104 ggca ctgatgtgat		20
<210>	105		
<211> <212>	18 DNA		
<213> <220>	Artificial Sequence		
<223>	Primer		
<400> gttggg	105 caga agagctag		18

<210>	106	
<211>	18	
<212>	DNA	
<213>	Artificial Sequence	
.000.		
<220> <223>	Primer	
(223/	rimei	
<400>	106	
	tagt ccccagag	18
,		
<210>	107	
<211>	21	
<212> <213>	DNA Artificial Sequence	
\213/	Artificial Sequence	
<220>		
<223>	Primer	
<400>	107	
tgcaag	agac acacatatga c	21
<210>	108	
<211>	21	
<212>	DNA	
<213>	Artificial Sequence	
<220>	Politica	
<223>	Primer	
<400>	108	
	ctgt tctcacaatt g	21
9		
<210>	109	
<211>	26	
<212> <213>	DNA Artificial Seguence	
(213)	Artificial Sequence	
<220>		
<223>	Primer	
<400>	109	
ggagcc	aagt ccatgtcact gaggga	26
<210>	110	
<211>	26	
<212>	DNA	
<213>	Artificial Sequence	
<220>		
<223>	Primer	
<400>	110	
	ttgc tgaggaacga aagaag	26
,	J J J	
<210>	111	
<211>	28	
<212> <213>	DNA Artificial Sequence	
\Z13/	Artificial Sequence	
<220>		
<223>	Primer	
<400>	111	~ ~
arccac	ttot toaacatact coacatoo	- 28

<210> <211> <212> <213>		
<220> <223>		
<400> gagagt	112 gaag aatgataagg acgcaaaa	28
<210><211><212><212><213>	DNA	
<220> <223>	Primer	
<400> ggtgtt	113 tttt tacactcttc cccccttgg	29
<210><211><211><212><213>	DNA	
<220> <223>	Primer	
<400> ccaagg	114 ccac aagattctcc c	21
<210><211><211><212><213>		
<220> <223>	Primer	
<400> agggga	115 gcca agtccatgtc acccagt	27
<210><211><211><212><213>	116 28 DNA Artificial Sequence	,
<220> <223>	Primer	
<400> gtttag	116 gact tgtttcggtt tggtggca	28
<210><211><211><212><213>	117 27 DNA Artificial Sequence	
<220> <223>	Primer	

```
<400> 117
tgccaccaaa ccgaaacaag tcctaaa
                                                                         27
<210> 118
<211> 26
<212> DNA
<213> Artificial Sequence
<220>
<223> Primer
<400> 118
catccacgga ccatcacttt ctgtta
                                                                         26
<210> 119
<211>
       28
<212>
       DNA
<213> Artificial Sequence
<220>
<223> Primer
<400> 119
tgaaatgaag ccaagtcctc aacatgag
                                                                         28
<210> 120
<211> 28
<212> DNA
<213> Artificial Sequence
<220>
<223> Primer
<400> 120
aaattacaga gagacaagct gccgctgc
                                                                         28
<210> 121
<211> 23
<212> PRT
<213> Artificial Sequence
<220>
<223> Synthetic Construct
<220>
<221> MISC FEATURE
<222>
       (1) ... (1)
<223> Xaa represents residues L, I, M, V or F
<220>
<221> misc_feature
<222>
      (2)..(3)
<223> Xaa can be any naturally occurring amino acid
<220>
<221> MISC_FEATURE
<222>
      (4)..(4)
<222> (4)..(4)
<223> Xaa represents residues L, I, M, V or F
<220>
<221> misc_feature
<222>
       (5)...(6)
<223> Xaa can be any naturally occurring amino acid
<220>
```

```
WARF0235.ST25.txt
<221> MISC_FEATURE
<222>
      (7)...(7)
<223> Xaa represents residues L, I, M, V or F
<220>
      misc_feature
<221>
<222>
      (8)..(9)
<223> Xaa can be any naturally occurring amino acid
<220>
<221> MISC_FEATURE
<222>
       (10)...(10)
      Xaa represents residues L, I, M, V or F
<223>
<220>
<221> misc feature
<222>
      (11)^{-}. (11)
<223> Xaa can be any naturally occurring amino acid
<220>
<221> MISC FEATURE
<222>
      (12)...(12)
<223> Xaa represents residues L, I, M, V or F
<220>
<221> misc feature
<222>
      (13)...(14)
<223> Xaa can be any naturally occurring amino acid
<220>
<221> MISC_FEATURE
<222>
       (15)...(15)
<223> Xaa represents residues N or C
<220>
<221>
      misc feature
<222>
      (16)^{-}. (17)
<223> Xaa can be any naturally occurring amino acid
<220>
<221> MISC_FEATURE
<222>
      (18)...(18)
<223> Xaa represents residues L, I, M, V or F
<220>
<221>
      misc feature
<222>
      (19)..(20)
<223> Xaa can be any naturally occurring amino acid
<220>
<221> MISC_FEATURE
<222>
      (21)^{-}.(21)
```

Xaa represents residues L, I, M, V or  ${\tt F}$ 

<223> Xaa can be any naturally occurring amino acid

10

Xaa Xaa Xaa Xaa Xaa Xaa

20

<210> 122

<223> <220>

<222>

<400> 121

<221> misc\_feature

(22)...(23)

15

```
<211> 12
<212> PRT
<213> Artificial Sequence
<220>
<223> Synthetic Construct
<220>
<221> MISC FEATURE
<222>
      (1)..(1)
<223> Xaa represents residues L, I, M, V or F
<220>
<221> misc_feature
<222>
      (2)..(3)
<223> Xaa can be any naturally occurring amino acid
<220>
<221> MISC_FEATURE
<222>
      (4)..(4)
<223> Xaa represents residues L, I, M, V or F
<220>
<221> misc_feature
<222>
      (5)...(6)
<223> Xaa can be any naturally occurring amino acid
<220>
<221> MISC FEATURE
<222>
      (7)...(7)
<223> Xaa represents residues L, I, M, V or F
<220>
<221> misc_feature
<222>
      (8)..(9)
<223> Xaa can be any naturally occurring amino acid
<220>
<221> MISC FEATURE
<222> (10)...(10)
<223> Xaa represents residues L, I, M, V or F
<220>
<221> misc_feature
<222>
      (11)^{-}. (11)
<223> Xaa can be any naturally occurring amino acid
<220>
<221> MISC FEATURE
<222>
      (12)...(12)
<223> Xaa represents residues L, I, M, V or F
<400> 122
<210> 123
<211> 6
<212> PRT
<213> Artificial Sequence
<220>
<223> Synthetic Construct
<220>
<221> MISC_FEATURE
```

```
WARF0235.ST25.txt
<222>
      (1)..(1)
<223> Xaa represents residues L, I, M, V or F
<220>
<221> misc_feature
<222>
       (2)..(3)
<223> Xaa can be any naturally occurring amino acid
<220>
<221> MISC FEATURE
<222>
       (4)...(4)
<223> Xaa represents residues L, I, M, V or F
<220>
<221> misc_feature
<222>
       (5)..(5)
<223> Xaa can be any naturally occurring amino acid
<220>
<221> MISC_FEATURE
<222>
       (6)...(6)
<223> Xaa represents residues L, I, M, V or F
<400> 123
Xaa Xaa Xaa Xaa Xaa
<210> 124
<211>
<212>
      PRT
<213> Artificial Sequence
<220>
<223>
      Synthetic Construct
<220>
<221> MISC FEATURE
<222>
      (1)...(1)
<223> Xaa represents residues L, I, M, V or F
<220>
<221> misc_feature
<222>
       (2)...(3)
<223> Xaa can be any naturally occurring amino acid
<220>
<221> MISC_FEATURE
<222>
      (4)..(4)
<223> Xaa represents residues L, I, M, V or F
<220>
<221>
      misc_feature
<222>
       (5)...(6)
<223> Xaa can be any naturally occurring amino acid
<220>
      MISC FEATURE
<221>
<222>
       (7)...(7)
<223> Xaa represents residues L, I, M, V or F
<220>
<221>
      misc feature
<222>
       (8)..(8)
<223>
      Xaa can be any naturally occurring amino acid
```

<220>

<221> MISC FEATURE

Page 95

```
WARF0235.ST25.txt
<222> (9)..(9) <223> Xaa represents residues L, I, M, V or F
<400> 124
Xaa Xaa Xaa Xaa Xaa Xaa Xaa
<210> 125
<211> 7
<212> PRT
<213> Artificial Sequence
<220>
<223> Synthetic Construct
<220>
<221> MISC_FEATURE
<222> (1)...(1) <223> Xaa represents residues L, I, M, V or F
<220>
<221> misc_feature
<222>
       (2) ... (3)
<223> Xaa can be any naturally occurring amino acid
<220>
<221> MISC FEATURE
<222> (4)..(4)
<223> Xaa represents residues L, I, M, V or F
<220>
<221> misc_feature
<222> (5)..(6)
<223> Xaa can be any naturally occurring amino acid
<220>
<221> MISC_FEATURE
<222> (7)...(7) <223> Xaa represents residues L, I, M, V or F
Xaa Xaa Xaa Xaa Xaa Xaa
```

Page 96